



## VA's Million Veteran Program Publications through May 2022

VA



U.S. Department of Veterans Affairs

Veterans Health Administration  
Office of Research & Development

Publication Title	Journal	Year	PMID/Journal
Million Veteran Program: A mega-biobank to study genetic influences on health and disease	Journal of Clinical Epidemiology	2016	<a href="#">26441289</a>
Exome-wide association study of plasma lipids in >300,000 individuals	Nature Genetics	2017	<a href="#">29083408</a>
Validating harmful alcohol use as a phenotype for genetic discovery using phosphatidylethanol and a polymorphism in ADH1B	Alcoholism Clinical and Experimental Research	2017	<a href="#">28295416</a>
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	<a href="#">JHRR</a>
Prevalence of Ideal Cardiovascular Health Metrics in the Million Veteran Program	American Journal of Cardiology	2018	<a href="#">29753396</a>
Alcohol Consumption and Risk of Coronary Artery Disease (From The Million Veteran Program)	American Journal of Cardiology	2018	<a href="#">29580627</a>
DASH Score and Subsequent Risk of Coronary Artery Disease: The Findings From Million Veteran Program	Journal of the American Heart Association	2018	<a href="#">29680824</a>
A phenotyping algorithm to identify acute ischemic stroke accurately from a national biobank: the Million Veteran Program	Clinical Epidemiology	2018	<a href="#">30425582</a>
Effects of Genetic Variants Associated with Familial Hypercholesterolemia on Low-Density Lipoprotein Cholesterol Levels and Cardiovascular Outcomes in the Million Veteran Program	Circulation Genomics and Precision Medicine	2018	<a href="#">31106297</a>
Genetics of blood lipids among ~300,000 multi-ethnic participants of the Million Veteran Program	Nature Genetics	2018	<a href="#">30275531</a>
Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits	Nature Genetics	2018	<a href="#">30224653</a>
Association of Interleukin 6 Receptor Variant With Cardiovascular Disease Effects of Interleukin 6 Receptor Blocking Therapy: A Phenome-Wide Association Study	JAMA Cardiology	2018	<a href="#">30090940</a>
Longitudinal drinking patterns and their clinical correlates in Million Veteran Program participants	Alcoholism Clinical and Experimental Research	2018	<a href="#">30592535</a>

AUDIT-C and ICD codes as phenotypes for harmful alcohol use: Association with ADH1B polymorphisms in two US populations	Addiction	2018	<a href="#">29972609</a>
Using DNA methylation to validate an electronic medical record phenotype for smoking	Addiction Biology	2018	<a href="#">30284751</a>
Genomics of Posttraumatic Stress Disorder in Veterans: Methods and Rationale for Veterans Affairs Cooperative Study #575B	International Journal of Methods in Psychiatric Research	2019	<a href="#">30767326</a>
Validation of an Electronic Medical Record-Based Algorithm for Identifying Posttraumatic Stress Disorder in US Veterans	Journal of Traumatic Stress	2019	<a href="#">31009556</a>
Genomewide Association Study of Maximum Habitual Alcohol Intake in >140,000 US European- and African-American Veterans Yields Novel Risk Loci	Biological Psychiatry	2019	<a href="#">31151762</a>
Genomewide Association Study of Posttraumatic Stress Disorder Re-Experiencing Symptoms in >165,000 US Veterans Yields New Biological Knowledge	Nature Neuroscience	2019	<a href="#">31358989</a>
International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci	Nature Communications	2019	<a href="#">31594949</a>
High-throughput Multimodal Automated Phenotyping (MAP) with Application to PheWAS	Journal of the American Medical Informatics Association	2019	<a href="#">JAMIA</a>
Gender Differences in Demographic and Health Characteristics of the Million Veteran Program Cohort	Women's Health Issues	2019	<a href="#">31253243</a>
Fried Food Consumption and Risk of Coronary Artery Disease: The Million Veteran Program	Clinical Nutrition	2019	<a href="#">31279615</a>
Omega-3 supplement use, fish intake, and risk of non-fatal coronary artery disease and	Clinical Nutrition	2019	<a href="#">30914216</a>
Association of APOL1 Risk Alleles with Cardiovascular Disease in African Americans in the Million Veteran Program	Circulation	2019	<a href="#">31337231</a>
Trans-ethnic association study of blood pressure determinants in over 750,000 individuals	Nature Genetics	2019	<a href="#">30578418</a>
Genome-wide association study of peripheral artery disease in the Million Veteran Program	Nature Medicine	2019	<a href="#">31285632</a>
Genomic and Transcriptomic Association Studies Identify 16 Novel Susceptibility Loci for Venous Thromboembolism	Blood	2019	<a href="#">31420334</a>
Harmonizing genetic ancestry and self-identified race/ethnicity in genome-wide association studies	American Journal of Human Genetics	2019	<a href="#">31564439</a>
Mapping eGFR loci to the renal transcriptome and phenome in the VA Million Veteran Program	Nature Communications	2019	<a href="#">31451708</a>

Genome-wide association analysis of venous thromboembolism identifies new risk loci and genetic overlap with arterial vascular disease	Nature Genetics	2019	<a href="#">5395047</a>
Genome-Wide Association Study of Apparent Treatment-Resistant Hypertension in the CHARGE Consortium: The CHARGE Pharmacogenetics Working Group	American Journal of Hypertension	2019	<a href="#">31545351</a>
Genome-wide association meta-analyses and fine-mapping elucidate pathways influencing albuminuria	Nature Communication	2019	<a href="#">31511532</a>
Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels	Nature Genetics	2019	<a href="#">31578528</a>
A catalog of genetic loci associated with kidney function from analyses of a million individuals	Nature Genetics	2019	<a href="#">31152163</a>
Genome-wide association study of alcohol consumption and use disorder in 274,424 individuals from multiple populations	Nature Communications	2019	<a href="#">30940813</a>
Genetically-guided algorithm development and sample size optimization for age-related macular degeneration cases and controls in electronic health records from the VA Million Veteran Program	AMIA Joint Summits on Translational Science Proceedings	2019	<a href="#">31258967</a>
Genome-Wide Association Studies of Schizophrenia and Bipolar Disorder in a Diverse Cohort of US Veterans	Schizophrenia Bulletin	2020	<a href="#">33169155</a>
Reproducible Genetic Risk Loci for Anxiety: Results From ~200,000 Participants in the Million Veteran Program	The American Journal of Psychiatry	2020	<a href="#">31906708</a>
Standardized Architecture for a Mega-Biobank Phenomic Library: The Million Veteran Program (MVP)	American Medical Informatics Association (AMIA) – Joint Summits on Translational Science proceedings	2020	<a href="#">32477652</a>
JSONize: A Scalable Machine Learning Pipeline to Model Medical Notes as Semi-structured Documents	American Medical Informatics Association (AMIA) – Joint Summits on Translational Science proceedings	2020	<a href="#">32477675</a>
PCSK9 loss of function is protective against extra-coronary atherosclerotic cardiovascular disease in a large multi-ethnic cohort	PLoS One	2020	<a href="#">33166319</a>
The relationship between circulating lipids and breast cancer risk: A Mendelian randomization study	PLoS Medicine	2020	<a href="#">32915777</a>
Egg consumption and risk of coronary artery disease in the Million Veteran Program	Clinical Nutrition	2020	<a href="#">31902601</a>
Genetic determinants of increased body mass index mediate the effect of smoking on increased risk for type 2 diabetes risk but not coronary artery disease	Human Molecular Genetics	2020	<a href="#">32833022</a>

Mendelian Randomization Analysis of Hemostatic Factors and Their Contribution to Peripheral Artery Disease	AHA Arteriosclerosis, Thrombosis, & Vascular Biology	2020	<a href="#">32847391</a>
Chromosome 1q21.2 and additional loci influence risk of spontaneous coronary artery dissection and myocardial infarction	Nature Communications	2020	<a href="#">32887874</a>
Genetic predisposition to smoking in relation to 14 cardiovascular diseases	European Heart Journal	2020	<a href="#">32300774</a>
Genetic Architecture of Abdominal Aortic Aneurysm in the Million Veteran Program	Circulation	2020	<a href="#">32981348</a>
Validating a non-invasive, ALT-based non-alcoholic fatty liver phenotype in the million veteran program	PLoS One	2020	<a href="#">32841307</a>
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	<a href="#">32541925</a>
Association of OPRM1 Functional Coding Variant With Opioid Use Disorder: A Genome-Wide Association Study	JAMA Psychiatry	2020	<a href="#">32492095</a>
Genome-wide association study of smoking trajectory and meta-analysis of smoking status in 842,000 individuals	Nature Communications	2020	<a href="#">33082346</a>
Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into biology and relationships with other traits	Nature Neuroscience	2020	<a href="#">32451486</a>
Novel Risk Loci in Tinnitus and Causal Inference With Neuropsychiatric Disorders Among Adults of European Ancestry	JAMA Otolaryngology - Head & Neck Surgery	2020	<a href="#">32970095</a>
Genotyping Array Design and Data Quality Control in the Million Veteran Program	American Journal of Human Genetics	2020	<a href="#">32243820</a>
Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population Architecture using Genomics and Epidemiology (PAGE) study	PLOS Genetics	2020	<a href="#">32226016</a>
A missense variant in Mitochondrial Amidoxime Reducing Component 1 gene and protection against liver disease	PLOS Genetics	2020	<a href="#">32282858</a>
The Polygenic and Monogenic Basis of Blood Traits and Diseases	Cell Press	2020	<a href="#">32888494</a>
Trans-ethnic and Ancestry-Specific Blood-Cell Genetics in 746,667 Individuals from 5 Global Populations	Cell Press	2020	<a href="#">32888493</a>
Genome-wide association analyses of post-traumatic stress disorder and its symptom subdomains in the Million Veteran Program	Nature Genetics	2021	<a href="#">33510476</a>

The Structure of Relationships between the Human Exposome and Cardiometabolic Health: The Million Veteran Program	Nutrients	2021	<a href="#">33921792</a>
Racial and Ethnic Disparities in U.S. Veteran Health Characteristics	International Journal of Environmental Research and Public Health	2021	<a href="#">33801200</a>
Consumption of potatoes and incidence rate of coronary artery disease: The Million Veteran Program	Clinical Nutrition ESPEN	2021	<a href="#">33745578</a>
Associations of biogeographic ancestry with hypertension traits	Journal of Hypertension	2021	<a href="#">33534346</a>
Urate, Blood Pressure, and Cardiovascular Disease: Evidence From Mendelian Randomization and Meta-Analysis of Clinical Trials	Hypertension	2021	<a href="#">33356394</a>
Genetics of Smoking and Risk of Atherosclerotic Cardiovascular Diseases: A Mendelian Randomization Study	JAMA Network Open	2021	<a href="#">33464320</a>
Multi-trait association studies discover pleiotropic loci between Alzheimer's disease and cardiometabolic traits	Alzheimer's Research & Therapy	2021	<a href="#">33541420</a>
Actionable druggable genome-wide Mendelian randomization identifies repurposing opportunities for COVID-19	Nature Medicine	2021	<a href="#">33837377</a>
Analysis of genetic and clinical factors associated with buprenorphine response	Drug Alcohol Depend	2021	<a href="#">34488071</a>
Chocolate consumption and risk of coronary artery disease: the Million Veteran Program	Am J Clin Nutr	2021	<a href="#">34483344</a>
Polygenic Breast Cancer Risk for Women Veterans in the Million Veteran Program	JCO Precis Oncol	2021	<a href="#">34381935</a>
Hearing loss and tinnitus: association studies for complex-hearing disorders in mouse and man	Hum Genet	2021	<a href="#">34318347</a>
Associations of Genetically Predicted Lp(a) (Lipoprotein [a]) Levels With Cardiovascular Traits in Individuals of European and African Ancestry	Circ Genom Precis Med	2021	<a href="#">34282949</a>
Discovery and prioritization of variants and genes for kidney function in >1.2 million individuals	Nature Communications	2021	<a href="#">34272381</a>
Regulatory variants in TCF7L2 are associated with thoracic aortic aneurysm	American Journal of Human Genetics	2021	<a href="#">34265237</a>
Cooperative Studies Program (CSP) #572: A Study of Serious Mental Illness in Veterans as a Pathway to personalized medicine in Schizophrenia and Bipolar Illness	Personalized Medicine in Psychiatry	2021	<a href="#">34222732</a>

Genomics of Gulf War Illness in U.S. Veterans Who Served during the 1990-1991 Persian Gulf War: Methods and Rationale for Veterans Affairs Cooperative Study #2006	Brain Sciences	2021	<a href="#">34202057</a>
Association of Apparent Treatment-Resistant Hypertension With Differential Risk of End-Stage Kidney Disease Across Racial Groups in the Million Veteran Program	Hypertension	2021	<a href="#">34148359</a>
Potential causal effect of posttraumatic stress disorder on alcohol use disorder and alcohol consumption in individuals of European descent: A Mendelian Randomization Study	Alcoholism, Clinical and Experimental Research	2021	<a href="#">34120358</a>
Psychological trauma and the genetic overlap between posttraumatic stress disorder and major depressive disorder	Psychological Medicine	2021	<a href="#">34085609</a>
Association of the transthyretin variant V122I with polyneuropathy among individuals of African ancestry	Scientific Reports	2021	<a href="#">34079032</a>
Metabolic Traits and Stroke Risk in Individuals of African Ancestry: Mendelian Randomization Analysis	Stroke	2021	<a href="#">34078102</a>
Bi-ancestral depression GWAS in the Million Veteran Program and meta-analysis in >1.2 million individuals highlight new therapeutic directions	Nature Neuroscience	2021	<a href="#">34045744</a>
Dietary yogurt is distinct from other dairy foods in its association with circulating lipid profile: Findings from the Million Veteran Program	Clinical Nutrition ESPEN	2021	<a href="#">34024555</a>
Phenome-wide association of 1809 phenotypes and COVID-19 disease progression in the Veterans Health Administration Million Veteran Program	PLoS One	2021	<a href="#">33984066</a>
Genetic analysis in European ancestry individuals identifies 517 loci associated with liver enzymes	Nature Communications	2021	<a href="#">33972514</a>
Association Between Genetic Variation in Blood Pressure and Increased Lifetime Risk of Peripheral Artery Disease	Arterioscler Thromb and Vasc Biol	2021	<a href="#">33853351</a>
COVID-19 Insights Partnership: Leveraging big data from the Department of Veterans Affairs and supercomputers at the Department of Energy under the public health authority	JAMIA	2021	<a href="#">33779710</a>
Attachment Style Moderates Polygenic Risk for Posttraumatic Stress in United States Military Veterans: Results From the National Health and Resilience in Veterans Study	Biological Psychiatry	2021	<a href="#">33276944</a>
Leisure time physical activity, sedentary behavior, and risk of cardiovascular disease and mortality among US Veterans	J Nov Physiother Phys Rehabil	2021	<a href="#">35098260</a>

Attachment Style Moderates Polygenic Risk for Incident Posttraumatic Stress in U.S. Military Veterans: A 7-Year, Nationally Representative, Prospective Cohort Study	Biol Psychiatry	2021	<a href="#">34955171</a>
Characterization of Million Veteran Program (MVP) enrollees with Comprehensive Traumatic Brain Injury Evaluation (CTBIE) data: An analysis of neurobehavioral symptoms	J Psychiatr Res	2021	<a href="#">34942434</a>
The power of genetic diversity in genome-wide association studies of lipids	Nature	2021	<a href="#">34887591</a>
Enhancing Discovery of Genetic Variants for Posttraumatic Stress Disorder Through Integration of Quantitative Phenotypes and Trauma Exposure Information	Biol Psychiatry	2021	<a href="#">34865855</a>
Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors	Biol Psychiatry	2021	<a href="#">34861974</a>
Trellis for efficient data and task management in the VA Million Veteran Program	Sci Rep	2021	<a href="#">34853358</a>
Development and Evaluation of a Rules-based Algorithm for Primary Open-Angle Glaucoma in the VA Million Veteran Program	Ophthalmic Epidemiol	2021	<a href="#">34822319</a>
Clinical knowledge extraction via sparse embedding regression (KESER) with multi-center large scale electronic health record data	NPJ Digit Med	2021	<a href="#">34707226</a>
A statistical quality assessment method for longitudinal observations in electronic health record data with an application to the VA million veteran program	BMC Med Inform Decis Mak	2021	<a href="#">34670548</a>
Use of the PsycheMERGE Network to Investigate the Association Between Depression Polygenic Scores and White Blood Cell Count	JAMA Psychiatry	2021	<a href="#">34668925</a>
Genetic liability for substance use associated with medical comorbidities in electronic health records of African- and European-ancestry individuals	Addict Biol	2021	<a href="#">34611967</a>
Disentangling sex differences in the shared genetic architecture of posttraumatic stress disorder, traumatic experiences, and social support with body size and composition	Neurobiol Stress	2021	<a href="#">34611531</a>
Association of Nut Consumption with Risk of Stroke and Cardiovascular Disease: The Million Veteran Program	Nutrients	2021	<a href="#">34578913</a>
A Missense Variant in the IL-6 Receptor and Protection From Peripheral Artery Disease	Circ Res	2021	<a href="#">34547901</a>
Analysis of genetic and clinical factors associated with buprenorphine response	Drug Alcohol Depend	2021	<a href="#">34488071</a>

Associations of Genetically Predicted Lp(a) (Lipoprotein [a]) Levels With Cardiovascular Traits in Individuals of European and African Ancestry	Circ Genom Precis Med	2021	<a href="#">34282949</a>
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Potential causal effect of posttraumatic stress disorder on alcohol use disorder and alcohol consumption in individuals of European descent: A Mendelian Randomization Study	Alcohol Clin Exp Res	2021	<a href="#">34120358</a>
Metabolic Traits and Stroke Risk in Individuals of African Ancestry: Mendelian Randomization Analysis	Stroke	2021	<a href="#">34078102</a>
A genome-wide association study of suicide attempts in the million veterans program identifies evidence of pan-ancestry and ancestry-specific risk loci	Mol Psychiatry	2022	<a href="#">35347246</a>
Lipid traits and type 2 diabetes risk in African ancestry individuals: A Mendelian Randomization study	EBioMedicine	2022	<a href="#">35325778</a>
Degree of Adherence to Based Diet and Total and Cause-Specific Mortality: Prospective Cohort Study in the Million Veteran Program	Public Health Nutr	2022	<a href="#">35307047</a>
Development of a polygenic risk score to improve detection of peripheral artery disease	Vascular Medicine	2022	<a href="#">35287516</a>
Dietary Sodium and Potassium Intake and Risk of Non-Fatal Cardiovascular Diseases: The Million Veteran Program	Nutrients	2022	<a href="#">35268096</a>
Genetically regulated multi-omics study for symptom clusters of posttraumatic stress disorder highlights pleiotropy with hematologic and cardio-metabolic traits	Molecular Psychiatry	2022	<a href="#">35241783</a>
Genetic and clinical determinants of abdominal aortic diameter: genome-wide association studies, exome array data and Mendelian randomization study	Human Molecular Genetics	2022	<a href="#">35234888</a>
Genetic Loci Associated With COVID-19 Positivity and Hospitalization in White, Black, and Hispanic Veterans of the VA Million Veteran Program	Frontiers in Genetics	2022	<a href="#">35222515</a>
Coronary Artery Disease Risk of Familial Hypercholesterolemia Genetic Variants Independent of Clinically Observed Longitudinal Cholesterol Exposure	Circulation Genomic and Precision Medicine	2022	<a href="#">35143253</a>
Transcriptome-wide association study in UK biobank Europeans identifies associations with blood cell traits	Human Molecular Genetics	2022	<a href="#">35138379</a>
Genome-wide meta-analysis of alcohol use disorder in East Asians	Neuropsychopharmacology	2022	<a href="#">35094024</a>



Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors	Biological Psychiatry	2022	<a href="#">34861974</a>
Genetic liability for substance use associated with medical comorbidities in electronic health records of African- and European-ancestry individuals	Addiction Biology	2022	<a href="#">34611967</a>
Genome-wide and phenome-wide analysis of ideal cardiovascular health in the VA Million Veteran Program	PLoS One	2022	<a href="#">35613103</a>
The Interrelationship of Tinnitus and Hearing Loss Secondary to Age, Noise Exposure, and Traumatic Brain Injury	Ear and hearing	2022	<a href="#">35612496</a>
Integration of rare expression outlier-associated variants improves polygenic risk prediction	American Journal of Human Genetics	2022	<a href="#">35588732</a>
A case for developing domain-specific vocabularies for extracting suicide factors from healthcare notes	Journal of Psychiatric Research	2022	<a href="#">35533516</a>
Subjective cognitive and psychiatric well-being in U.S. Military Veterans screened for deployment-related traumatic brain injury: A Million Veteran Program Study	Journal of Psychiatric Research	2022	<a href="#">35483132</a>
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	<a href="#">35482673</a>
Million Veteran Program's response to COVID-19: Survey development and preliminary findings	PLoS One	2022	<a href="#">35468170</a>
Integrating human brain proteomes with genome-wide association data implicates novel proteins in post-traumatic stress disorder	Molecular Psychiatry	2022	<a href="#">35449297</a>
Binary acronym disambiguation in clinical notes from electronic health records with an application in computational phenotyping	Internal journal of medical informatics	2022	<a href="#">35405530</a>
A genome-wide association study of suicide attempts in the million veterans program identifies evidence of pan-ancestry and ancestry-specific risk loci	Molecular Psychiatry	2022	<a href="#">35347246</a>
Coronary Artery Disease Risk of Familial Hypercholesterolemia Genetic Variants Independent of Clinically Observed Longitudinal Cholesterol Exposure	Circular, Genomic and precision medicine	2022	<a href="#">35143253</a>
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	<a href="#">35089317</a>
Attachment Style Moderates Polygenic Risk for Incident Posttraumatic Stress in U.S. Military Veterans: A 7-Year, Nationally Representative, Prospective Cohort Study	Biological Psychiatry	2022	<a href="#">34955171</a>

Enhancing Discovery of Genetic Variants for Posttraumatic Stress Disorder Through Integration of Quantitative Phenotypes and Trauma Exposure Information	Biological Psychiatry	2022	<a href="#">34865855</a>
Hearing loss and tinnitus: association studies for complex-hearing disorders in mouse and man	Human Genetics	2022	<a href="#">34318347</a>