

VA's Million Veteran Program Publications through December 2021



U.S. Department of Veterans Affairs

Publication Title	Journal	Year	PMID/Journal
Genome-wide association analyses of post-traumatic stress disorder and its symptom subdomains in the Million Veteran Program	Nature Genetics	2021	<u>33510476</u>
The Structure of Relationships between the Human Exposome and Cardiometabolic Health: The Million Veteran Program	Nutrients	2021	<u>33921792</u>
Racial and Ethnic Disparities in U.S. Veteran Health Characteristics	International Journal of Environmental Research and Public Health	2021	<u>33801200</u>
Consumption of potatoes and incidence rate of coronary artery disease: The Million Veteran Program	Clinical Nutrition ESPEN	2021	<u>33745578</u>
Associations of biogeographic ancestry with hypertension traits	Journal of Hypertension	2021	<u>33534346</u>
Urate, Blood Pressure, and Cardiovascular Disease: Evidence From Mendelian Randomization and Meta-Analysis of Clinical Trials	Hypertension	2021	<u>33356394</u>
Genetics of Smoking and Risk of Atherosclerotic Cardiovascular Diseases: A Mendelian Randomization Study	JAMA Network Open	2021	<u>33464320</u>
Multi-trait association studies discover pleiotropic loci between Alzheimer's disease and cardiometabolic traits	Alzheimer's Research & Therapy	2021	<u>33541420</u>
Actionable druggable genome-wide Mendelian randomization identifies repurposing opportunities for COVID-19	Nature Medicine	2021	<u>33837377</u>
Analysis of genetic and clinical factors associated with buprenorphine response	Drug Alcohol Depend	2021	<u>34488071</u>
Chocolate consumption and risk of coronary artery disease: the Million Veteran Program	Am J Clin Nutr	2021	<u>34483344</u>
Polygenic Breast Cancer Risk for Women Veterans in the Million Veteran Program	JCO Precis Oncol	2021	<u>34381935</u>

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Hearing loss and tinnitus: association studies for complex-hearing disorders in mouse and man	Hum Genet	2021	<u>34318347</u>
Associations of Genetically Predicted Lp(a) (Lipoprotein [a]) Levels With Cardiovascular Traits in Individuals of European and African Ancestry	Circ Genom Precis Med	2021	<u>34282949</u>
Discovery and prioritization of variants and genes for kidney function in >1.2 million individuals	Nature Communications	2021	<u>34272381</u>
Regulatory variants in TCF7L2 are associated with thoracic aortic aneurysm	American Journal of Human Genetics	2021	<u>34265237</u>
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	<u>34222732</u>
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	<u>34202057</u>
Association of Apparent Treatment-Resistant Hypertension with Differential Risk of End- Stage Kidney Disease Across Racial Groups in the Million Veteran Program	Hypertension	2021	<u>34148359</u>
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	<u>34120358</u>
Psychological trauma and the genetic overlap between posttraumatic stress disorder and major depressive disorder	Psychological Medicine	2021	<u>34085609</u>
Association of the transthyretin variant V122I with polyneuropathy among individuals of African ancestry	Scientific Reports	2021	<u>34079032</u>
Metabolic Traits and Stroke Risk in Individuals of African Ancestry: Mendelian Randomization Analysis	Stroke	2021	<u>34078102</u>
Bi-ancestral depression GWAS in the Million Veteran Program and meta-analysis in >1.2 million individuals highlight new therapeutic directions	Nature Neuroscience	2021	<u>34045744</u>
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	<u>34024555</u>

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Phenome-wide association of 1809 phenotypes and COVID-19 disease progression in the Veterans Health Administration Million Veteran Program	PLoS One	2021	<u>33984066</u>
Genetic analysis in European ancestry individuals identifies 517 loci associated with liver enzymes	Nature Communications	2021	<u>33972514</u>
Association Between Genetic Variation in Blood Pressure and Increased Lifetime Risk of Peripheral Artery Disease	Arterioscler Thromb and Vasc Biol	2021	<u>33853351</u>
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	<u>33779710</u>
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	<u>33276944</u>
Leisure time physical activity, sedentary behavior, and risk of cardiovascular disease and mortality among US Veterans	J Nov Physiother Phys Rehabil	2021	<u>35098260</u>
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	<u>34955171</u>
Characterization of Million Veteran Program (MVP) enrollees with Comprehensive Traumatic Brain Injury Evaluation (CTBIE) data: An analysis of neurobehavioral symptoms	J Psychiatr Res	2021	<u>34942434</u>
The power of genetic diversity in genome-wide association studies of lipids	Nature	2021	<u>34887591</u>
Enhancing Discovery of Genetic Variants for Posttraumatic Stress Disorder Through Integration of Quantitative Phenotypes and Trauma Exposure Information	Biol Psychiatry	2021	<u>34865855</u>
Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors	Biol Psychiatry	2021	<u>34861974</u>
Trellis for efficient data and task management in the VA Million Veteran Program	Sci Rep	2021	<u>34853358</u>
Development and Evaluation of a Rules-based Algorithm for Primary Open-Angle Glaucoma in the VA Million Veteran Program	Ophthalmic Epidemiol	2021	<u>34822319</u>

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Clinical knowledge extraction via sparse embedding regression (KESER) with multi-center large scale electronic health record data	NPJ Digit Med	2021	<u>34707226</u>
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	<u>34670548</u>
Use of the PsycheMERGE Network to Investigate the Association Between Depression Polygenic Scores and White Blood Cell Count	JAMA Psychiatry	2021	<u>34668925</u>
Genetic liability for substance use associated with medical comorbidities in electronic health records of African- and European-ancestry individuals	Addict Biol	2021	<u>34611967</u>
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	<u>34611531</u>
Association of Nut Consumption with Risk of Stroke and Cardiovascular Disease: The Million Veteran Program	Nutrients	2021	<u>34578913</u>
A Missense Variant in the IL-6 Receptor and Protection from Peripheral Artery Disease	Circ Res	2021	<u>34547901</u>
Analysis of genetic and clinical factors associated with buprenorphine response	Drug Alcohol Depend	2021	<u>34488071</u>
Associations of Genetically Predicted Lp(a) (Lipoprotein [a]) Levels With Cardiovascular Traits in Individuals of European and African Ancestry	Circ Genom Precis Med	2021	<u>34282949</u>
Association of Apparent Treatment-Resistant Hypertension with Differential Risk of End- Stage Kidney Disease Across Racial Groups in the Million Veteran Program	Hypertension	2021	<u>34148359</u>
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	<u>34120358</u>
Metabolic Traits and Stroke Risk in Individuals of African Ancestry: Mendelian Randomization Analysis	Stroke	2021	<u>34078102</u>

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Genome-Wide Association Studies of Schizophrenia and Bipolar Disorder in a Diverse Cohort of US Veterans	Schizophrenia Bulletin	2020	<u>33169155</u>
Reproducible Genetic Risk Loci for Anxiety: Results From ~200,000 Participants in the Million Veteran Program	The American Journal of Psychiatry	2020	<u>31906708</u>
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	<u>32477652</u>
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	<u>32477675</u>
PCSK9 loss of function is protective against extra-coronary atherosclerotic cardiovascular disease in a large multi-ethnic cohort	PLoS One	2020	<u>33166319</u>
The relationship between circulating lipids and breast cancer risk: A Mendelian randomization study	PLoS Medicine	2020	<u>32915777</u>
Egg consumption and risk of coronary artery disease in the Million Veteran Program	Clinical Nutrition	2020	<u>31902601</u>
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	<u>32833022</u>
Mendelian Randomization Analysis of Hemostatic Factors and Their Contribution to Peripheral Artery Disease	AHA Arteriosclerosis, Thrombosis, & Vascular Biology	2020	<u>32847391</u>
Chromosome 1q21.2 and additional loci influence risk of spontaneous coronary artery dissection and myocardial infarction	Nature Communications	2020	<u>32887874</u>
Genetic predisposition to smoking in relation to 14 cardiovascular diseases	European Heart Journal	2020	<u>32300774</u>
Genetic Architecture of Abdominal Aortic Aneurysm in the Million Veteran Program	Circulation	2020	<u>32981348</u>

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Validating a non-invasive, ALT-based non-alcoholic fatty liver phenotype in the Million Veteran Program	PLoS One	2020	<u>32841307</u>
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	<u>32541925</u>
Association of OPRM1 Functional Coding Variant with Opioid Use Disorder: A Genome- Wide Association Study	JAMA Psychiatry	2020	<u>32492095</u>
Genome-wide association study of smoking trajectory and meta-analysis of smoking status in 842,000 individuals	Nature Communications	2020	<u>33082346</u>
Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into biology and relationships with other traits	Nature Neuroscience	2020	<u>32451486</u>
Novel Risk Loci in Tinnitus and Causal Inference with Neuropsychiatric Disorders Among Adults of European Ancestry	JAMA Otolaryngology - Head & Neck Surgery	2020	<u>32970095</u>
Genotyping Array Design and Data Quality Control in the Million Veteran Program	American Journal of Human Genetics	2020	<u>32243820</u>
Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population Architecture using Genomics and Epidemiology (PAGE) study	PLOS Genetics	2020	<u>32226016</u>
A missense variant in Mitochondrial Amidoxime Reducing Component 1 gene and protection against liver disease	PLOS Genetics	2020	<u>32282858</u>
The Polygenic and Monogenic Basis of Blood Traits and Diseases	Cell Press	2020	<u>32888494</u>
Trans-ethnic and Ancestry-Specific Blood-Cell Genetics in 746,667 Individuals from 5 Global Populations	Cell Press	2020	<u>32888493</u>
Genomics of Posttraumatic Stress Disorder in Veterans: Methods and Rationale for Veterans Affairs Cooperative Study #575B	International Journal of Methods in Psychiatric Research	2019	<u>30767326</u>
Validation of an Electronic Medical Record-Based Algorithm for Identifying Posttraumatic Stress Disorder in US Veterans	Journal of Traumatic Stress	2019	<u>31009556</u>
Genomewide Association Study of Maximum Habitual Alcohol Intake in >140,000 US European- and African-American Veterans Yields Novel Risk Loci	Biological Psychiatry	2019	<u>31151762</u>

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Genomewide Association Study of Posttraumatic Stress Disorder Re-Experiencing Symptoms in >165,000 US Veterans Yields New Biological Knowledge	Nature Neuroscience	2019	<u>31358989</u>
International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci	Nature Communications	2019	<u>31594949</u>
High-throughput Multimodal Automated Phenotyping (MAP) with Application to PheWAS	Journal of the American Medical Informatics Association	2019	<u>31613361</u>
Gender Differences in Demographic and Health Characteristics of the Million Veteran Program Cohort	Women's Health Issues	2019	<u>31253243</u>
Fried Food Consumption and Risk of Coronary Artery Disease: The Million Veteran Program	Clinical Nutrition	2019	<u>31279615</u>
Omega-3 supplement use, fish intake, and risk of non-fatal coronary artery disease and	Clinical Nutrition	2019	<u>30914216</u>
Association of APOL1 Risk Alleles with Cardiovascular Disease in African Americans in the Million Veteran Program	Circulation	2019	<u>31337231</u>
Trans-ethnic association study of blood pressure determinants in over 750,000 individuals	Nature Genetics	2019	<u>30578418</u>
Genome-wide association study of peripheral artery disease in the Million Veteran Program	Nature Medicine	2019	<u>31285632</u>
Genomic and Transcriptomic Association Studies Identify 16 Novel Susceptibility Loci for Venous Thromboembolism	Blood	2019	<u>31420334</u>
Harmonizing genetic ancestry and self-identified race/ethnicity in genome-wide association studies	American Journal of Human Genetics	2019	<u>31564439</u>
Mapping eGFR loci to the renal transcriptome and phenome in the VA Million Veteran Program	Nature Communications	2019	<u>31451708</u>
Genome-wide association analysis of venous thromboembolism identifies new risk loci and genetic overlap with arterial vascular disease	Nature Genetics	2019	<u>5395047</u>
Genome-Wide Association Study of Apparent Treatment-Resistant Hypertension in the CHARGE Consortium: The CHARGE Pharmacogenetics Working Group	American Journal of Hypertension	2019	<u>31545351</u>
Genome-wide association meta-analyses and fine-mapping elucidate pathways influencing albuminuria	Nature Communication	2019	<u>31511532</u>

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Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels	Nature Genetics	2019	<u>31578528</u>
A catalog of genetic loci associated with kidney function from analyses of a million individuals	Nature Genetics	2019	<u>31152163</u>
Genome-wide association study of alcohol consumption and use disorder in 274,424 individuals from multiple populations	Nature Communications	2019	<u>30940813</u>
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	<u>31258967</u>
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	<u>33117892</u>
Prevalence of Ideal Cardiovascular Health Metrics in the Million Veteran Program	American Journal of Cardiology	2018	<u>29753396</u>
Alcohol Consumption and Risk of Coronary Artery Disease (From the Million Veteran Program)	American Journal of Cardiology	2018	<u>29580627</u>
DASH Score and Subsequent Risk of Coronary Artery Disease: The Findings From Million Veteran Program	Journal of the American Heart Association	2018	<u>29680824</u>
A phenotyping algorithm to identify acute ischemic stroke accurately from a national biobank: the Million Veteran Program	Clinical Epidemiology	2018	<u>30425582</u>
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	<u>31106297</u>
Genetics of blood lipids among ~300,000 multi-ethnic participants of the Million Veteran Program	Nature Genetics	2018	<u>30275531</u>
Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits	Nature Genetics	2018	<u>30224653</u>
Association of Interleukin 6 Receptor Variant with Cardiovascular Disease Effects of Interleukin 6 Receptor Blocking Therapy: A Phenome-Wide Association Study	JAMA Cardiology	2018	<u>30090940</u>

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Longitudinal drinking patterns and their clinical correlates in Million Veteran Program participants	Alcoholism Clinical and Experimental Research	2018	<u>30592535</u>
AUDIT-C and ICD codes as phenotypes for harmful alcohol use: Association with ADH1B polymorphisms in two US populations	Addiction	2018	<u>29972609</u>
Using DNA methylation to validate an electronic medical record phenotype for smoking	Addiction Biology	2018	<u>30284751</u>
Exome-wide association study of plasma lipids in >300,000 individuals	Nature Genetics	2017	<u>29083408</u>
Validating harmful alcohol use as a phenotype for genetic discovery using phosphatidylethanol and a polymorphism in ADH1B	Alcoholism Clinical and Experimental Research	2017	<u>28295416</u>
Million Veteran Program: A mega-biobank to study genetic influences on health and disease	Journal of Clinical Epidemiology	2016	<u>26441289</u>