

Resource Summary Report

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Million Veteran Program

RRID:SCR_021731

Type: Tool

Proper Citation

Million Veteran Program (RRID:SCR_021731)

Resource Information

URL: <https://www.research.va.gov/mvp/>

Proper Citation: Million Veteran Program (RRID:SCR_021731)

Description: National research program to learn how genes, lifestyle, and military exposures affect health and illness.

Abbreviations: MVP

Resource Type: data or information resource, portal, topical portal

Keywords: Genomic dataset, genes, lifestyle, military exposures, affect health, affect illness

Availability: Free, Freely available

Resource Name: Million Veteran Program

Resource ID: SCR_021731

Record Creation Time: 20220129T080357+0000

Record Last Update: 20260821T194204+0000

Ratings and Alerts

No rating or validation information has been found for Million Veteran Program.

No alerts have been found for Million Veteran Program.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 273 mentions in open access literature.

Listed below are recent publications. The full list is available at [dkNET](#) .

Grotzinger AD, et al. (2026) Mapping the genetic landscape across 14 psychiatric disorders. *Nature*, 649(8096), 406.

Tong J, et al. (2026) Metabolic Improvement Mediates the Causal Relationship Between GLP-1 Receptor Agonists and Myocardial Infarction: A Mendelian Randomization and Mediation Analysis Study. *Diabetes care*, 49(1), 171.

Clark SL, et al. (2026) A large-scale DNA methylation study of alcohol use identified robust associations and cell-type specific insights. *Molecular psychiatry*, 31(6), 3506.

Insawang B, et al. (2026) Genome-wide association study of corneal dystrophy uncovers novel risk loci and enables improved polygenic prediction of Fuchs endothelial corneal dystrophy. *medRxiv : the preprint server for health sciences*.

Wu D, et al. (2026) Circulating Inflammatory Proteins and Fuchs Endothelial Corneal Dystrophy: A Mendelian Randomization and Bioinformatics-Based Investigation. *Translational vision science & technology*, 15(5), 6.

Topiwala A, et al. (2026) Alcohol use and risk of dementia in diverse populations: evidence from cohort, case-control and Mendelian randomisation approaches. *BMJ evidence-based medicine*, 31(1), 13.

Broberg M, et al. (2026) Genome-wide meta-analysis identifies genetic risk loci for mono- and polyneuropathies in 983 477 individuals. *Human molecular genetics*, 35(4).

Meng F, et al. (2026) Multi-layered functional genomics prioritizes candidate effectors and regulatory mechanisms of ankylosing spondylitis. *Frontiers in immunology*, 17, 1859310.

Wendland ZD, et al. (2026) A Genome-Wide Association Study of Hidradenitis Suppurativa from the VA's Million Veteran Program. *Dermatology (Basel, Switzerland)*, 242(4), 418.

Song JO, et al. (2026) African-enriched SLC39A10 (ZIP10) missense variants differentially affect cellular zinc homeostasis and associate with health-related traits. *Genes & nutrition*, 21(1).

Wang Y, et al. (2026) Biological age acceleration, genetic susceptibility, and incident diabetic retinopathy risk: A prospective cohort study. *The journal of nutrition, health & aging*, 30(6), 100863.

Kinzy TG, et al. (2026) A Primary Open-Angle Glaucoma Locus Near Transcription Factor PRRX1 Identified in the Million Veteran Program. *Ophthalmology science*, 6(7), 101196.

White DL, et al. (2026) Association between deployment and Gulf War Illness and adverse COVID outcomes in a nationwide cohort of 1990-1991 Gulf Era war veterans in the VA's Million Veteran Program. *PloS one*, 21(6), e0348594.

Barr PB, et al. (2026) Polygenic risk factors for comorbid diagnoses in individuals with substance use disorders: A phenome-wide survival analysis. *Psychological medicine*, 56, e174.

Huang L, et al. (2026) Association of SARS-CoV-2 infection with incident diabetes among U.S. Veterans in a prospective longitudinal cohort. *PloS one*, 21(6), e0351992.

Gasperi M, et al. (2026) A multi-ancestry meta genome-wide association study of migraine among veterans: associations with traumatic brain injury, depression, and post-traumatic stress disorder. *Molecular psychiatry*, 31(5), 2660.

Kamiza AB, et al. (2026) KidneyGenAfrica multi-cohort Genome-wide association study and polygenic prediction of kidney function in 110,000 Africans. *Nature communications*, 17(1).

Shi Z, et al. (2026) Genome-wide analysis implicates inner ear development in Ménière's disease. *medRxiv : the preprint server for health sciences*.

Davis CN, et al. (2026) Education shares distinct genetic influences with substance use and disorder. *Psychological medicine*, 56, e38.

Wang Y, et al. (2026) Causal Association Between Commonly Used Medicines and Diabetes-Related Eye Diseases: Univariable and Multivariable Mendelian Randomization Study. *Journal of diabetes research*, 2026(1), e6075025.