

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, topical portal, 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: 20260811T043654+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 [RRID](#) .

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.

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.

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.

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.