

## MVP Publications – April 2021



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Veterans Health Administration  
Office of Research & Development

| Associated MVP Project or Core Team | Publication Title  | First Author(s)         | Last Author(s)  | Journal  | Year | PMID/Journal             |
|-------------------------------------|--|-------------------------|-----------------|--|------|--------------------------|
| CSP572, MVP000                      | Genome-Wide Association Studies of Schizophrenia and Bipolar Disorder in a Diverse Cohort of US Veterans                                       | Bigdeli, Tim            | Harvey, Philip  | Schizophrenia Bulletin                                   | 2020 | <a href="#">33169155</a> |
| CSP575b                             | Genomics of Posttraumatic Stress Disorder in Veterans: Methods and Rationale for Veterans Affairs Cooperative Study #575B                      | Radhakrishnan, Krishnan | Gelernter, Joel | International Journal of Methods in Psychiatric Research | 2019 | <a href="#">30767326</a> |
| CSP575b                             | Validation of an Electronic Medical Record-Based Algorithm for Identifying Posttraumatic Stress Disorder in US Veterans                        | Harrington, Kelly       | Cho, Kelly      | Journal of Traumatic Stress                              | 2019 | <a href="#">31009556</a> |
| CSP575b                             | Genomewide Association Study of Maximum Habitual Alcohol Intake in >140,000 US European- and African-American Veterans Yields Novel Risk Loci  | Gelernter, Joel         | Stein, Murray   | Biological Psychiatry                                    | 2019 | <a href="#">31151762</a> |
| CSP575b                             | Genomewide Association Study of Posttraumatic Stress Disorder Re-Experiencing Symptoms in >165,000 US Veterans Yields New Biological Knowledge | Gelernter, Joel         | Stein, Murray   | Nature Neuroscience                                      | 2019 | <a href="#">31358989</a> |
| CSP575b                             | International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci                    | Nievergelt, CM          | Koenen, KC      | Nature Communications                                    | 2019 | <a href="#">31594949</a> |
| CSP575b                             | Reproducible Genetic Risk Loci for Anxiety: Results From ~200,000 Participants in the Million Veteran Program                                  | Levey, Daniel           | Stein, Murray   | The American Journal of Psychiatry                       | 2020 | <a href="#">31906708</a> |
| CSP575b                             | Genome-wide association analyses of post-traumatic stress disorder and its symptom subdomains in the Million Veteran Program                   | Stein, Murray           | Gelernter, Joel | Nature Genetics  | 2021 | <a href="#">33510476</a> |

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| MVP000                              | The Structure of Relationships between the Human Exposome and Cardiometabolic Health: The Million Veteran Program | Ivey, Kerry L.    | Djousse, Luc          | Nutrients  | 2021 | <a href="#">33921792</a> |
| MVP000                              | Racial and Ethnic Disparities in U.S. Veteran Health Characteristics  | Ward, Rachel E.   | Whitbourne, Stacey B. | International Journal of Environmental Research and Public Health                | 2021 | <a href="#">33801200</a> |
| MVP000                              | High-throughput Multimodal Automated Phenotyping (MAP) with Application to PheWAS                                 | Liao, Katherine   | Cai, Tianxi           | Journal of the American Medical Informatics Association                          | 2019 | <a href="#">JAMIA</a>    |
| MVP000                              | Baseline Characterization and Annual Trends of Body Mass Index for a Mega Biobank Cohort of US Veterans 2011-2017 | Nguyen, Xuan-Mai  | Cho, Kelly            | Journal of Health Research and Reviews   | 2018 | <a href="#">JHRR</a>     |
| MVP000                              | Prevalence of Ideal Cardiovascular Health Metrics in the Million Veteran Program                                  | Nguyen, Xuan-Mai  | Djousse, Luc          | American Journal of Cardiology   | 2018 | <a href="#">29753396</a> |
| MVP000                              | Million Veteran Program: A mega-biobank to study genetic influences on health and disease                         | Gaziano, Michael  | O'Leary, Timothy      | Journal of Clinical Epidemiology   | 2016 | <a href="#">26441289</a> |
| MVP000                              | Gender Differences in Demographic and Health Characteristics of the Million Veteran Program Cohort                | Harrington, Kelly | Whitbourne, Stacey    | Women's Health Issues  | 2019 | <a href="#">31253243</a> |
| MVP000, CHAMPION                    | Standardized Architecture for a Mega-Biobank Phenomic Library: The Million Veteran Program (MVP)                  | Knight, Kathryn E | Cho, Kelly            | American Medical Informatics Association (AMIA) – Joint Summits on Translational | 2020 | <a href="#">32477652</a> |

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|                                     |   |                                  |                  | Science proceedings  |      |                          |
| MVP000, CHAMPION                    | JSONize: A Scalable Machine Learning Pipeline to Model Medical Notes as Semi-structured Documents                                 | Rush, Everett                    | Begoli, Edmon    | American Medical Informatics Association (AMIA) – Joint Summits on Translational Science proceedings | 2020 | <a href="#">32477675</a> |
| MVP000, MVP001                      | Consumption of potatoes and incidence rate of coronary artery disease: The Million Veteran Program                                | Bodar, Vijaykumar                | Djousse, Luc     | Clinical Nutrition ESPEN   | 2021 | <a href="#">33745578</a> |
| MVP000, MVP001                      | Fried Food Consumption and Risk of Coronary Artery Disease: The Million Veteran Program   | Honerlaw, Jacqueline             | Djousse, Luc     | Clinical Nutrition   | 2019 | <a href="#">31279615</a> |
| MVP000, MVP001                      | Alcohol Consumption and Risk of Coronary Artery Disease (From The Million Veteran Program)  | Song, Rebecca                    | Djousse, Luc     | American Journal of Cardiology   | 2018 | <a href="#">29580627</a> |
| MVP000, MVP001                      | Omega-3 supplement use, fish intake, and risk of non-fatal coronary artery disease and  | Ward, Rachel                     | Djousse, Luc     | Clinical Nutrition   | 2019 | <a href="#">30914216</a> |
| MVP000, MVP001                      | DASH Score and Subsequent Risk of Coronary Artery Disease: The Findings From Million Veteran Program                              | Djousse, Luc                     | Gaziano, Michael | Journal of the American Heart Association  | 2018 | <a href="#">29680824</a> |
| MVP000, MVP001, MVP003, MVP028      | PCSK9 loss of function is protective against extra-coronary atherosclerotic cardiovascular disease in a large multi-ethnic cohort | Small, Aeron                     | Damrauer, Scott  | PLoS One   | 2020 | <a href="#">33166319</a> |
| MVP000, MVP003                      | The relationship between circulating lipids and breast cancer risk: A Mendelian randomization study                               | Johnson, Kelsey<br>Klarin, Derek | Voight, Derek    | PLoS Medicine  | 2020 | <a href="#">32915777</a> |

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| MVP001                              | Egg consumption and risk of coronary artery disease in the Million Veteran Program   | Djousse, Luc  | Cho, Kelly  | Clinical Nutrition                          | 2020 | <a href="#">31902601</a> |
| MVP001                              | A phenotyping algorithm to identify acute ischemic stroke accurately from a national biobank: the Million Veteran Program  | Imran, Tasnim   | Cho, Kelly  | Clinical Epidemiology                       | 2018 | <a href="#">30425582</a> |
| MVP001, MVP002, MVP003              | Association of APOL1 Risk Alleles with Cardiovascular Disease in African Americans in the Million Veteran Program  | Bick, Alexander<br>Akwo, Elvis<br>Robinson-Cohen,<br>Cassieanne<br>Lee, K | Hung, Adriana<br>Damrauer, Scott  | Circulation                                 | 2019 | <a href="#">31337231</a> |
| MVP001, MVP002, MVP003              | Trans-ethnic association study of blood pressure determinants in over 750,000 individuals  | Giri, Ayush<br>Hellwege, Jacklyn<br>Keaton, Jacob<br>Park, Jihwan         | Susztak, Katalin<br>O'Donnell,<br>Christopher<br>Hung, Adriana<br>Edwards, Todd   | Nature Genetics                             | 2019 | <a href="#">30578418</a> |
| MVP001, MVP003                      | Genome-wide association study of peripheral artery disease in the Million Veteran Program  | Klarin, Derek   | Damrauer, Scott<br>Tsao, Philip   | Nature Medicine                             | 2019 | <a href="#">31285632</a> |
| MVP001, MVP003                      | Effects of Genetic Variants Associated with Familial Hypercholesterolemia on Low-Density Lipoprotein Cholesterol Levels and Cardiovascular Outcomes in the Million Veteran Program | Sun, Yan  | Wilson, Peter   | Circulation Genomics and Precision Medicine | 2018 | <a href="#">31106297</a> |
| MVP001, MVP003                      | Genetics of blood lipids among ~300,000 multi-ethnic participants of the Million Veteran Program   | Klarin, Derek<br>Damrauer, Scott  | Assimes,<br>Themistocles<br>Wilson, Peter<br>Rader, Daniel<br>Kathiresan,<br>Sekar<br>Tsao, Philip<br>O'Donnell,<br>Christopher | Nature Genetics                             | 2018 | <a href="#">30275531</a> |

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| MVP001, MVP003                      | Genomic and Transcriptomic Association Studies Identify 16 Novel Susceptibility Loci for Venous Thromboembolism  | Lindstrom, S   | Smith, NL   | Blood                              | 2019 | <a href="#">31420334</a> |
| MVP001, MVP003                      | Harmonizing genetic ancestry and self-identified race/ethnicity in genome-wide association studies   | Fang, H  | Tang, H   | American Journal of Human Genetics | 2019 | <a href="#">31564439</a> |
| MVP001, MVP003                      | 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     | Thom, Christopher S.   | Voight, Benjamin F.   | Human Molecular Genetics           | 2020 | <a href="#">32833022</a> |
| MVP001, MVP002, MVP003              | Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits  | Evangelou, Evangelos<br>Warren, Helen<br>Mosen-Ansorena, David<br>Mifsud, Borbala<br>Pazoki, Raha<br>Gao, He<br>Ntritsos, Georgios<br>Dimou, Niki<br>Tzoulaki, Ioanna<br>Barnes, Michael<br>Wain, Louise<br>Elliott, Paul<br>Caulfield, Mark | Hung, Adriana<br>O'Donnell, Christopher<br>Edwards, Todd<br>Psaty, Bruce<br>Tzoulaki, Ioanna<br>Barnes, Michael<br>Wain, Louise<br>Elliott, Paul<br>Caulfield, Mark | Nature Genetics                    | 2018 | <a href="#">30224653</a> |
| MVP001, MVP002, MVP003              | Mapping eGFR loci to the renal transcriptome and phenome in the VA Million Veteran Program   | Hellwege, Jacklyn<br>Velez Edwards, Digna<br>Giri, Ayush   | Susztak, Katalin<br>Edwards, Todd<br>Hung, Adriana  | Nature Communications              | 2019 | <a href="#">31451708</a> |
| MVP001, MVP003                      | Association of Interleukin 6 Receptor Variant With Cardiovascular Disease Effects of Interleukin 6 Receptor Blocking Therapy: A Phenome-Wide Association Study | Cai, Tianxi<br>Zhang, Yichi  | Liao, Katherine   | JAMA Cardiology                    | 2018 | <a href="#">30090940</a> |

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| MVP001, MVP003                      | Genome-wide association analysis of venous thromboembolism identifies new risk loci and genetic overlap with arterial vascular disease         | Klarin, Derek  | Natarajan, Pradeep<br>Damrauer, Scott  | Nature Genetics                                      | 2019 | <a href="#">5395047</a>  |
| MVP001, MVP003                      | Exome-wide association study of plasma lipids in >300,000 individuals  | Liu, Dajiang,<br>Peloso, Gina<br>Yu, Haojie<br>Butterworth, Adam<br>Wang, Xiao<br>Mahajan, Anubha<br>Saleheen, Danish<br>Emdin, Connor<br>Willer, Cristen<br>Kathiresan, Sekar | Howson, Joanna<br>Danesh, John<br>McCarthy, Mark<br>Cowan, Chad<br>Abecasis, Goncalo<br>Deloukas, Panos<br>Musunuru, Kiran<br>Willer, Cristen<br>Kathiresan, Sekar | Nature Genetics                                      | 2017 | <a href="#">29083408</a> |
| MVP001, MVP003                      | Mendelian Randomization Analysis of Hemostatic Factors and Their Contribution to Peripheral Artery Disease                                     | Small, Aeron<br>Huffman, Jennifer E  | Damrauer, Scott  | AHA Arteriosclerosis, Thrombosis, & Vascular Biology | 2020 | <a href="#">32847391</a> |
| MVP001, MVP003                      | Chromosome 1q21.2 and additional loci influence risk of spontaneous coronary artery dissection and myocardial infarction                       | Saw, Jacqueline  | Ganesh, Santhi   | Nature Communications                                | 2020 | <a href="#">32887874</a> |
| MVP001, MVP002, MVP027              | Associations of biogeographic ancestry with hypertension traits  | Keaton, Jacob M.   | Velez Edwards, Digna R.  | Journal of Hypertension                              | 2021 | <a href="#">33534346</a> |
| MVP002                              | Genome-Wide Association Study of Apparent Treatment-Resistant Hypertension in the CHARGE Consortium: The CHARGE Pharmacogenetics Working Group | Irvin, MR  | Ahluwalia, TS<br>Arnett, AK<br>Avery, CL   | American Journal of Hypertension                     | 2019 | <a href="#">31545351</a> |
| MVP002                              | Genome-wide association meta-analyses and fine-mapping elucidate pathways  | Teumer, Alexander<br>Li, Yong<br>Ghasemi, Sahar  | Butterworth, Adam<br>Hung, Adriana   | Nature Communication                                 | 2019 | <a href="#">31511532</a> |

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|                                     | influencing albuminuria   | Prins, Bram<br>Wuttke, Matthias<br>Hermle, Tobias  | Pattaro, Cristian<br>Kottgen, Anna  |                                |      |                          |
| MVP002                              | Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels                             | Tin, A<br>Marten, J<br>Halperin Kuhns, VL<br>Li, Y<br>Wuttke, M<br>Kirsten, H                  | Hung, AM<br>Teumer, A<br>Pattaro, C<br>Woodward, OM<br>Vitart, V<br>Köttgen, A          | Nature Genetics                | 2019 | <a href="#">31578528</a> |
| MVP002                              | A catalog of genetic loci associated with kidney function from analyses of a million individuals                              | Wuttke, Matthias<br>Li, Yong<br>Li, Man<br>Sieber, Karsten<br>Feitosa, Mary<br>Gorski, Mathias | Heid, Iris<br>Scholz, Markus<br>Teumer, Alexander<br>Kottgen, Anna<br>Pattaro, Cristian | Nature Genetics                | 2019 | <a href="#">31152163</a> |
| MVP003                              | Genetic predisposition to smoking in relation to 14 cardiovascular diseases   | Larsson, Susanna   | Burgess, Stephen  | European Heart Journal         | 2020 | <a href="#">32300774</a> |
| MVP003                              | Urate, Blood Pressure, and Cardiovascular Disease: Evidence From Mendelian Randomization and Meta-Analysis of Clinical Trials | Gill, Dipender   | Tzoulaki, Ioanna  | Hypertension                   | 2021 | <a href="#">33356394</a> |
| MVP003                              | Genetics of Smoking and Risk of Atherosclerotic Cardiovascular Diseases: A Mendelian Randomization Study                      | Levin, Michael   | Voight, Benjamin  | JAMA Network Open              | 2021 | <a href="#">33464320</a> |
| MVP003<br>MVP000                    | Genetic Architecture of Abdominal Aortic Aneurysm in the Million Veteran Program  | Klarin, Derek  | Tsao, Philip  | Circulation                    | 2020 | <a href="#">32981348</a> |
| MVP003<br>MVP004                    | Genome-wide association study of alcohol consumption and use disorder in 274,424 individuals from multiple populations        | Kranzler, Henry<br>Zhou, Hang<br>Kember, Rachel  | Gelernter, Joel   | Nature Communications          | 2019 | <a href="#">30940813</a> |
| MVP003,<br>MVP028                   | Multi-trait association studies discover pleiotropic loci between Alzheimer's disease and cardiometabolic traits              | Bone, William P.   | Voight, Benjamin F.   | Alzheimer's Research & Therapy | 2021 | <a href="#">33541420</a> |

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| MVP003, MVP028                      | Validating a non-invasive, ALT-based non-alcoholic fatty liver phenotype in the million veteran program   | Serper, Marina   | Chang, Kyong-Mi  | PLoS One                                      | 2020 | <a href="#">32841307</a> |
| MVP003 MVP028                       | 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 | Vujkovic, Marijana   | Saleheen, Danish | Nature Genetics                               | 2020 | <a href="#">32541925</a> |
| MVP004                              | Longitudinal drinking patterns and their clinical correlates in Million Veteran Program participants  | Vickers Smith, Rachel<br>Kranzler, Henry<br>Justice, Amy   | Tate, Janet      | Alcoholism Clinical and Experimental Research | 2018 | <a href="#">30592535</a> |
| MVP004                              | AUDIT-C and ICD codes as phenotypes for harmful alcohol use: Association with ADH1B polymorphisms in two US populations                           | Justice, Amy<br>Smith, Rachel<br>Tate, Janet<br>McGinnis, Kathleen<br>Xu, Ke<br>Becker, William<br>Lee, Kuang-Yao<br>Lynch, Kevin<br>Sun, Ning<br>Concato, John<br>Fiellin, David<br>Zhao, Hongyu<br>Gelernter, Joel | Kranzler, Henry  | Addiction                                     | 2018 | <a href="#">29972609</a> |
| MVP004                              | Using DNA methylation to validate an electronic medical record phenotype for smoking  | McGinnis, Kathleen<br>Justice, Amy<br>Tate, Janet<br>Kranzler, Henry<br>Tindle, Hiliary<br>Becker, William<br>Concato, John<br>Gelernter, Joel<br>Li, Boyang<br>Zhang Xinyu  | Xu, Ke           | Addiction Biology                             | 2018 | <a href="#">30284751</a> |



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|                                     |  | Zhao, Hongyu<br>Crothers, Kristina  |                         |   |      |                          |
| MVP004                              | Validating harmful alcohol use as a phenotype for genetic discovery using phosphatidylethanol and a polymorphism in ADH1B  | Justice, Amy<br>McGinnis, Kathleen<br>Tate, Janet<br>Xu, Ke<br>Becker, William<br>Zhao, Hongyu<br>Gelernter, Joel | Kranzler, Henry         | Alcoholism<br>Clinical and<br>Experimental<br>Research              | 2017 | <a href="#">28295416</a> |
| MVP004                              | Association of OPRM1 Functional Coding Variant With Opioid Use Disorder: A Genome-Wide Association Study   | Zhou, Hang  | Gelernter, Joel         | JAMA<br>Psychiatry  | 2020 | <a href="#">32492095</a> |
| MVP004                              | Genome-wide association study of smoking trajectory and meta-analysis of smoking status in 842,000 individuals   | Xu, Ke<br>Li, Boyang  | Justice, Amy            | Nature<br>Communications  | 2020 | <a href="#">33082346</a> |
| MVP004<br>CSP575b<br>MVP025         | Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into biology and relationships with other traits   | Zhou, Hang  | Gelernter, Joel         | Nature<br>Neuroscience  | 2020 | <a href="#">32451486</a> |
| MVP005                              | 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 | Halladay, Chris   | Crawford, Dana          | AMIA Joint<br>Summits on<br>Translational<br>Science<br>Proceedings | 2019 | <a href="#">31258967</a> |
| MVP010                              | Novel Risk Loci in Tinnitus and Causal Inference With Neuropsychiatric Disorders Among Adults of European Ancestry   | Clifford, Royce   | Nievergelt,<br>Caroline | JAMA<br>Otolaryngology -<br>Head & Neck<br>Surgery                  | 2020 | <a href="#">32970095</a> |
| MVP035                              | Actionable druggable genome-wide Mendelian randomization identifies repurposing opportunities for COVID-19   | Gaziano, Liam   | Casas, Juan P.          | Nature Medicine   | 2021 | <a href="#">33837377</a> |

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| MVP Genomic Working Group           | Genotyping Array Design and Data Quality Control in the Million Veteran Program  | Hunter-Zinck, Haley | Pyarajan, Saiju   | American Journal of Human Genetics | 2020 | <a href="#">32243820</a> |
| Collaboration                       | Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population Architecture using Genomics and Epidemiology (PAGE) study | Hu, Yao             | Peters, Ulrike    | PLOS Genetics                      | 2020 | <a href="#">32226016</a> |
| Collaboration                       | A missense variant in Mitochondrial Amidoxime Reducing Component 1 gene and protection against liver disease   | Emdin, Connor       | Kathiresan, Sekar | PLOS Genetics                      | 2020 | <a href="#">32282858</a> |
| Collaboration                       | The Polygenic and Monogenic Basis of Blood Traits and Diseases   | Vuckovic, Dragana   | Soranzo, Nicole   | Cell Press                         | 2020 | <a href="#">32888494</a> |
| Collaboration                       | Trans-ethnic and Ancestry-Specific Blood-Cell Genetics in 746,667 Individuals from 5 Global Populations  | Chen, Ming-Huei     | Lettre, Guillaume | Cell Press                         | 2020 | <a href="#">32888493</a> |