Resource Summary Report

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University of Pennsylvania Perelman School of Medicine Penn Medicine BioBank Core Facility

RRID:SCR_022415
Type: Tool

Proper Citation

University of Pennsylvania Perelman School of Medicine Penn Medicine BioBank Core Facility (RRID:SCR_022415)

Resource Information

URL: https://pmbb.med.upenn.edu/

Proper Citation: University of Pennsylvania Perelman School of Medicine Penn Medicine BioBank Core Facility (RRID:SCR_022415)

Description: BioBank supports researchers by providing centralized access to large number of annotated blood and tissue samples.

Abbreviations: PMBB

Synonyms: University of Pennsylvania Perelman School of Medicine Penn Medicine BioBank

Resource Type: core facility, storage service resource, service resource, access service resource, material storage repository, biobank

Keywords: USEDit, ABRF, annotated blood and tissue samples

Resource Name: University of Pennsylvania Perelman School of Medicine Penn Medicine BioBank Core Facility

Resource ID: SCR_022415

Alternate IDs: ARBF_1421

Alternate URLs: https://coremarketplace.org?citation=1&FacilityID=1421

Record Creation Time: 20220602T050140+0000

Record Last Update: 20260819T044540+0000

Ratings and Alerts

No rating or validation information has been found for University of Pennsylvania Perelman School of Medicine Penn Medicine BioBank Core Facility.

No alerts have been found for University of Pennsylvania Perelman School of Medicine Penn Medicine BioBank Core Facility.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 16 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF .

Levin MG, et al. (2026) Genetic Prediction of Circulating Lipoprotein(a) Levels in Diverse Populations. medRxiv : the preprint server for health sciences.

Hartmann K, et al. (2026) Association of a polygenic risk score with coronary atherosclerotic burden in clinical CT angiograms. medRxiv : the preprint server for health sciences.

Li C, et al. (2026) Proteomic risk score for early prediction of kidney disease progression in individuals with APOL1 high-risk genotypes. Nature medicine, 32(5), 1701.

Cousins KAQ, et al. (2026) Electronic Health Records to Test Multimorbidity Influences to Plasma Biomarker Interpretation for Alzheimer's Disease. Annals of neurology, 99(4), 1030.

Cousins KAQ, et al. (2025) Electronic health records to test multimorbidity influences to plasma biomarker interpretation for Alzheimer's disease. medRxiv : the preprint server for health sciences.

Hui D, et al. (2025) Risk factors affecting polygenic score performance across diverse cohorts. eLife, 12.

DePaolo J, et al. (2025) Titin-Truncating variants predispose to dilated cardiomyopathy in populations genetically similar to african and european reference populations. PLoS genetics, 21(6), e1011727.

Nicastro OR, et al. (2025) Demographic and Genetic Predictors of ADHD Diagnoses and Medications in Electronic Health Records. Research square.

Bao MM, et al. (2025) Phenome-wide association study of monogenic inflammatory bowel disease genes in diverse biobanks identifies population-specific and shared Goldilocks alleles: implications for Precision Medicine. Journal of Crohn's & colitis, 19(7).

DePaolo J, et al. (2025) Genome-First Approach to Rare and Common Variant Risk of Thoracic Aortic Aneurysm and Dissection. medRxiv : the preprint server for health

sciences.

Corbett RJ, et al. (2025) Germline pathogenic variation impacts somatic alterations and patient outcomes in pediatric central nervous system tumors. Nature communications, 16(1), 10282.

Neylan CJ, et al. (2025) Novel polygenic risk score associates with diverticulitis in a multi-institutional, ancestrally diverse cohort. Scientific reports, 15(1), 42348.

Zhang DY, et al. (2024) Protein-truncating variant in APOL3 increases chronic kidney disease risk in epistasis with APOL1 risk alleles. JCI insight, 9(19).

DePaolo J, et al. (2024) Titin-Truncating variants Predispose to Dilated Cardiomyopathy in Diverse Populations. medRxiv : the preprint server for health sciences.

Cappadocia J, et al. (2024) PMS2CL interference leading to erroneous identification of a pathogenic PMS2 variant in Black patients. Genetics in medicine open, 2, 101858.

Klarin D, et al. (2023) Genome-wide association study of thoracic aortic aneurysm and dissection in the Million Veteran Program. Nature genetics, 55(7), 1106.