

Resource Summary Report

Generated by RRID on Aug 4, 2026

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

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: 20260804T043756+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 10 mentions in open access literature.

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

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.

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

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).

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.

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

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.