Resource Summary Report

Generated by SPARC SAWG on Sep 19, 2026

All of Us

RRID:SCR_027032
Type: Tool

Proper Citation

All of Us (RRID:SCR_027032)

Resource Information

URL: https://www.researchallofus.org

Proper Citation: All of Us (RRID:SCR_027032)

Description: Portal stores health data from participants from across the United States. Provides interactive Data Browser where anyone can learn about the type and quantity of data that All of Us collects. Users can explore aggregate data including genomic variants, survey responses, physical measurements, electronic health record information, and wearables data.

Resource Type: data or information resource, database, portal

Keywords: health data, United States data, genomic variants, survey responses, physical measurements, electronic health record information,

Availability: Free, Freely available,

Resource Name: All of Us

Resource ID: SCR_027032

Record Creation Time: 20250605T060221+0000

Record Last Update: 20260912T200517+0000

Ratings and Alerts

No rating or validation information has been found for All of Us.

No alerts have been found for All of Us.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 82 mentions in open access literature.

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

Ak??imen F, et al. (2026) Long-read sequencing identifies FGF14 repeat expansions in Parkinson's disease. Brain : a journal of neurology, 149(5), 1514.

Jang SK, et al. (2026) Leveraging protein language models to identify complex trait associations with previously inaccessible classes of functional rare variants. Cell genomics, 6(2), 101068.

Wu E, et al. (2026) Neoplasms Associated with the Onset of Age-Related Macular Degeneration: A Case-Control Study Using the All of Us Cohort. Research square.

Zhang Y, et al. (2026) Shared genetic architecture between sleep traits and cardiometabolic diseases. EBioMedicine, 126, 106220.

Hwang I, et al. (2026) Association of GLP-1 Receptor Agonists With Hepatic Decompensation in the All of Us Research Program. Diabetes, obesity & metabolism, 28(6), 4837.

Fulda ES, et al. (2026) 11 million days of longitudinal wearable data reveal novel future health insights. medRxiv : the preprint server for health sciences.

Nyeo SS, et al. (2026) Population-scale sequencing resolves determinants of persistent EBV DNA. Nature, 650(8102), 664.

Chan ZM, et al. (2026) Heart rate phase is an indicator of chronotype-relevant circadian shifts associated with human disease: an All of Us Research Program analysis. The Journal of physiology, 604(8), 3282.

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

Arizpe A, et al. (2026) Disparities in multidimensional psychosocial stressors by sexual identity among cancer survivors from the All of Us Research Program. Cancer causes & control : CCC, 37(5).

De Jonghe J, et al. (2026) Saturation editing of RNU4-2 reveals distinct dominant and recessive disorders. Nature, 654(8118), 429.

Zamora G, et al. (2026) Comparing random forest and elastic net models to predict substance use disorder transitions in participants with cannabis and stimulant use: Evidence from the All of Us cohort. Drug and alcohol dependence, 278, 113012.

Baierl J, et al. (2026) S4-multi: Enhancing polygenic score prediction in ancestrally diverse populations. HGG advances, 7(1), 100551.

Vand K, et al. (2026) Genomic Disaggregation Reveals Distinct Admixture Patterns and Cardiometabolic Risk Loci in Black Hawaiians. bioRxiv : the preprint server for biology.

Wang Y, et al. (2026) Phenome-derived polygenic scores and social determinants jointly shape context-dependent disease risk. medRxiv : the preprint server for health sciences.

Shi Z, et al. (2026) CalPred yields calibrated intervals for polygenic risk prediction. medRxiv : the preprint server for health sciences.

Long X, et al. (2026) Linkage between HLA-B8 and HLA-DQ2.5 Contributes to Ancestry-Dependent Genetic Risk for Celiac Disease. medRxiv : the preprint server for health sciences.

Winters AH, et al. (2026) A Genome-First Study of Familial Hypercholesterolemia Comparing African and European Ancestry Individuals. Circulation, 153(24), 1928.

Yang E, et al. (2026) Racial and Ethnic Disparities in Degenerative Lumbar Disc Disease: A Population-Based Study Using the All of Us Research Program. Global spine journal, 16(5), 2272.

Cruz-Vespa N, et al. (2026) Depression Risk in Type 1 Versus Type 2 Diabetes: Cross-Sectional Analysis of Body Mass Index (BMI) in a Nationally Diverse Cohort. Endocrinology, diabetes & metabolism, 9(2), e70172.