Resource Summary Report

Generated by RRID on Sep 10, 2026

Analysis, Visualization, and Informatics Lab-space (AnVIL)

RRID:SCR_017469
Type: Tool

Proper Citation

Analysis, Visualization, and Informatics Lab-space (AnVIL) (RRID:SCR_017469)

Resource Information

URL: https://anvilproject.org/

Proper Citation: Analysis, Visualization, and Informatics Lab-space (AnVIL) (RRID:SCR_017469)

Description: Portal to facilitate integration and computing on and across large datasets generated by NHGRI programs, as well as initiatives funded by National Institutes of Health or by other agencies that support human genomics research. Resource for genomic scientific community, that leverages cloud based infrastructure for democratizing genomic data access, sharing and computing across large genomic, and genomic related data sets. Component of federated data ecosystem, and is expected to collaborate and integrate with other genomic data resources through adoption of FAIR (Findable, Accessible, Interoperable, Reusable) principles, as their specifications emerge from scientific community. Will provide collaborative environment, where datasets and analysis workflows can be shared within consortium and be prepared for public release to broad scientific community through AnVIL user interfaces.

Abbreviations: AnVIL

Synonyms: Visualization, and Informatics Lab-space, AnVIL, Analysis Visualization and Informatics Lab-space, Analysis

Resource Type: data or information resource, data repository, portal, project portal, service resource, storage service resource

Keywords: Dataset, NHGRI, program, NIH, initiative, funded, human, genomic, data, access, sharing, FAIR, analysis, workflow

Funding: NIH

Availability: Restricted

Resource Name: Analysis, Visualization, and Informatics Lab-space (AnVIL)

Resource ID: SCR_017469

Alternate URLs: https://www.genome.gov/Funded-Programs-Projects/Computational-Genomics-and-Data-Science-Program/Genomic-Analysis-Visualization-Informatics-Lab-space-AnVIL

Record Creation Time: 20220129T080335+0000

Record Last Update: 20260905T132824+0000

Ratings and Alerts

No rating or validation information has been found for Analysis, Visualization, and Informatics Lab-space (AnVIL).

No alerts have been found for Analysis, Visualization, and Informatics Lab-space (AnVIL).

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 31 mentions in open access literature.

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

Calame DG, et al. (2026) PTPN1-related autoinflammation is a common cause of Aicardi-Goutières Syndrome with reduced penetrance. medRxiv : the preprint server for health sciences.

Meyer-Schuman R, et al. (2026) A massively parallel reporter assay of MECP2 cis-regulatory elements reveals genetic candidates for male-biased autism. bioRxiv : the preprint server for biology.

Pehlivan D, et al. (2026) Bi-allelic variants in NRDC cause a neurodevelopmental disorder characterized by neonatal lethality, microcephaly, and brain abnormalities. American journal of human genetics, 113(3), 548.

Isaac KJ, et al. (2025) Insights into the Datasets, Tools, and Training Needs of the AnVIL Community: 2024. bioRxiv : the preprint server for biology.

Manolio TA, et al. (2025) Advancing the science of genomic learning healthcare systems. Learning health systems, 9(4), e70027.

Dardas Z, et al. (2025) Bi-allelic UGGT1 variants cause a congenital disorder of glycosylation. American journal of human genetics, 112(5), 1139.

Palou-Márquez G, et al. (2025) Variable efficiency of nonsense-mediated mRNA decay across human tissues, tumors and individuals. Genome biology, 26(1), 316.

Speir ML, et al. (2025) Making genomic data FAIR through effective Data Portals. Scientific data, 12(1), 1872.

Genner R, et al. (2024) Assessing methylation detection for primary human tissue using Nanopore sequencing. bioRxiv : the preprint server for biology.

Iyer S, et al. (2024) The BRAIN Initiative data-sharing ecosystem: Characteristics, challenges, benefits, and opportunities. eLife, 13.

Wang Z, et al. (2024) NCI Cancer Research Data Commons: Resources to Share Key Cancer Data. Cancer research, 84(9), 1388.

Hou K, et al. (2024) Admix-kit: an integrated toolkit and pipeline for genetic analyses of admixed populations. Bioinformatics (Oxford, England), 40(4).

Ng JK, et al. (2024) HAT: de novo variant calling for highly accurate short-read and long-read sequencing data. Bioinformatics (Oxford, England), 40(1).

Guitart X, et al. (2024) Independent expansion, selection and hypervariability of the TBC1D3 gene family in humans. bioRxiv : the preprint server for biology.

Pitsava G, et al. (2024) Genome sequencing reveals the impact of non-canonical exon inclusions in rare genetic disease. medRxiv : the preprint server for health sciences.

Nagasaki M, et al. (2023) Design and implementation of a hybrid cloud system for large-scale human genomic research. Human genome variation, 10(1), 6.

Seddighi S, et al. (2023) Mis-spliced transcripts generate de novo proteins in TDP-43-related ALS/FTD. bioRxiv : the preprint server for biology.

Liao WW, et al. (2023) A draft human pangenome reference. Nature, 617(7960), 312.

Hou K, et al. (2023) Admix-kit: An Integrated Toolkit and Pipeline for Genetic Analyses of Admixed Populations. bioRxiv : the preprint server for biology.

Shih CC, et al. (2023) A five-safes approach to a secure and scalable genomics data repository. iScience, 26(4), 106546.