

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 7, 2026

Genomic Datasharing

RRID:SCR_005233

Type: Tool

Proper Citation

Genomic Datasharing (RRID:SCR_005233)

Resource Information

URL: <http://gds.nih.gov/>

Proper Citation: Genomic Datasharing (RRID:SCR_005233)

Description: NIH established expectations for sharing data obtained through NIH-funded genome-wide association studies (GWAS) with the implementation of the GWAS Policy. Information and resources related to the GWAS Policy can be found on this website.

Abbreviations: GDS

Synonyms: GWAS, Genome-Wide Association Studies

Resource Type: data or information resource, feed, listserv, narrative resource, standard specification

Keywords: genome-wide association study, genome, data sharing

Funding: NIH

Resource Name: Genomic Datasharing

Resource ID: SCR_005233

Alternate IDs: OMICS_00272

Record Creation Time: 20220129T080229+0000

Record Last Update: 20260905T132532+0000

Ratings and Alerts

No rating or validation information has been found for Genomic Datasharing.

No alerts have been found for Genomic Datasharing.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Falk MJ, et al. (2015) Mitochondrial Disease Sequence Data Resource (MSeqDR): a global grass-roots consortium to facilitate deposition, curation, annotation, and integrated analysis of genomic data for the mitochondrial disease clinical and research communities. *Molecular genetics and metabolism*, 114(3), 388.

Lehner T, et al. (2015) Convergence of advances in genomics, team science, and repositories as drivers of progress in psychiatric genomics. *Biological psychiatry*, 77(1), 6.

Frank E, et al. (2015) All the world's a (clinical) stage: rethinking bipolar disorder from a longitudinal perspective. *Molecular psychiatry*, 20(1), 23.

Berger S, et al. (2015) Effectiveness of shrinkage and variable selection methods for the prediction of complex human traits using data from distantly related individuals. *Annals of human genetics*, 79(2), 122.

Paltoo DN, et al. (2014) Data use under the NIH GWAS data sharing policy and future directions. *Nature genetics*, 46(9), 934.