

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

caGWAS

RRID:SCR_009617

Type: Tool

Proper Citation

caGWAS (RRID:SCR_009617)

Resource Information

URL: <https://wiki.nci.nih.gov/display/caGWAS/caGWAS>

Proper Citation: caGWAS (RRID:SCR_009617)

Description: Tool that allows researchers to integrate, query, report, and analyze significant associations between genetic variations and disease, drug response or other clinical outcomes. SNP array technologies make it possible to genotype hundreds of thousands of single nucleotide polymorphisms (SNPs) simultaneously, enabling whole genome association studies. Within the Clinical Genomic Object Model (CGOM), the caIntegrator team created a domain model for Whole Genome Association Study Analysis. CGOM-caGWAS is a semantically annotated domain model that captures associations between Study, Study Participant, Disease, SNP Association Analysis, SNP Population Frequency and SNP annotations. caGWAS APIs and web portal provide: * a semantically annotated domain model, database schema with sample data, seasoned middleware, APIs, and web portal for GWAS data; * platform and disease agnostic CGOM-caGWAS model and associated APIs; * the opportunity for developers to customize the look and feel of their GWAS portal; * a foundation of open source technologies; * a well-tested and performance-enhanced platform, as the same software is being used to house the CGEMS data portal; * accelerated analysis of results from various biomedical studies; and * a single application through which researchers and bioinformaticians can access and analyze clinical and experimental data from a variety of data types, as caGWAS objects are part of the CGOM, which includes microarray, genomic, immunohistochemistry, imaging, and clinical data.

Abbreviations: caGWAS

Synonyms: caGWAS (Cancer Genome-Wide Association Studies), Cancer Genome-Wide Association Studies, CGOM-caGWAS

Resource Type: software resource

Keywords: application, computational neuroscience, genetic association, genomic analysis, imaging genomics, java, snp, gene, software, web environment, microarray,

genomic, immunohistochemistry, imaging, clinical

Availability: BSD License

Resource Name: caGWAS

Resource ID: SCR_009617

Alternate IDs: nlx_155841

Alternate URLs: <http://www.nitrc.org/projects/cagwas>

Record Creation Time: 20220129T080254+0000

Record Last Update: 20260905T132635+0000

Ratings and Alerts

No rating or validation information has been found for caGWAS.

No alerts have been found for caGWAS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.