GWAS Central

RRID:SCR_006170
Type: Tool

Proper Citation

GWAS Central (RRID:SCR_006170)

Resource Information

**URL:** [http://www.gwascentral.org/](http://www.gwascentral.org/)

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**Description:** Publicly available database of summary level findings from genetic association studies in humans, including genome wide association studies (GWAS). Previously named HGBASE, HGVbase and HGVbaseG2P.

**Synonyms:** Human Genome Variation database of Genotype to Phenotype information, HGVbaseG2P, Genome Wide Association Studies Central

**Resource Type:** storage service resource, database, service resource, data or information resource, data repository

**Defining Citation:** PMID:18948288

**Keywords:** Human Genome Variation database of Genotype-to-Phenotype information, genetic association study, genotype, phenotype, gene, genome region, disease, frequency data, region, genome, marker, single nucleotide polymorphism, genetic variant, allele, genome wide association study, human genome, chromosome, genetics

**Funding Agency:** European Union GEN2PHEN project, University of Leicester; Leicester; United Kingdom, GlaxoSmithKline

**Resource Name:** GWAS Central

**Resource ID:** SCR_006170

**Alternate IDs:** nlx_151672, nif-0000-02958, nif-0000-02958, SCR_007709
Alternate URLs: http://www.hgvbaseg2p.org

Ratings and Alerts

No rating or validation information has been found for GWAS Central.

No alerts have been found for GWAS Central.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 69 mentions in open access literature.

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


Schulze KV, et al. (2019) Edematous severe acute malnutrition is characterized by hypomethylation of DNA. Nature communications, 10(1), 5791.