Resource Summary Report

Generated by SPARC SAWG on Sep 18, 2026

GeneCruiser

RRID:SCR_003153
Type: Tool

Proper Citation

GeneCruiser (RRID:SCR_003153)

Resource Information

URL: http://genecruiser.broadinstitute.org/genecruiser3/

Proper Citation: GeneCruiser (RRID:SCR_003153)

Description: A web service and web application for the annotation of microarray data providing integrated access to genomic information freely available from public data sources.

Abbreviations: GeneCruiser

Resource Type: data access protocol, service resource, software resource, web service

Defining Citation: PMID:16030072

Keywords: gene, genetic variation, probe, variation, annotation

Availability: Free, Freely available

Resource Name: GeneCruiser

Resource ID: SCR_003153

Alternate IDs: OMICS_00760

Alternate URLs: https://www.broadinstitute.org/publications/broad3691

Record Creation Time: 20220129T080217+0000

Record Last Update: 20260912T195552+0000

Ratings and Alerts

No rating or validation information has been found for GeneCruiser.

No alerts have been found for GeneCruiser.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Robiou-du-Pont S, et al. (2015) Should we have blind faith in bioinformatics software? Illustrations from the SNAP web-based tool. PloS one, 10(3), e0118925.

Vaughan LK, et al. (2015) Linkage and association analysis of obesity traits reveals novel loci and interactions with dietary n-3 fatty acids in an Alaska Native (Yup'ik) population. Metabolism: clinical and experimental, 64(6), 689.

Faino A, et al. (2014) Identifying rare variants associated with hypertension using the C-alpha test. BMC proceedings, 8(Suppl 1 Genetic Analysis Workshop 18Vanessa Olmo), S56.

Vaughan LK, et al. (2013) Where in the genome are we? A cautionary tale of database use in genomics research. Frontiers in genetics, 4, 38.