

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 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:** 20260905T132458+0000

### Ratings and Alerts

No rating or validation information has been found for GeneCruiser.

No alerts have been found for GeneCruiser.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 4 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [Kravitz](#) .

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.