Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

GENSENG

RRID:SCR_000378
Type: Tool

Proper Citation

GENSENG (RRID:SCR_000378)

Resource Information

URL: http://sourceforge.net/projects/genseng/

Proper Citation: GENSENG (RRID:SCR_000378)

Description: Software for detecting copy number variations from next generation sequencing data. Used to identify regions of discrete copy number changes while simultaneously accounting for effects of multiple confounders.

Resource Type: data analysis software, data processing software, software application, software resource

Defining Citation: PMID:23275535

Keywords: next generation sequencing data, detecting copy number variations, discrete copy number changes, identify regions

Availability: Free, Available for download, Freely available

Resource Name: GENSENG

Resource ID: SCR_000378

Alternate IDs: OMICS_00345

Record Creation Time: 20220129T080201+0000

Record Last Update: 20260905T132418+0000

Ratings and Alerts

No rating or validation information has been found for GENSENG.

No alerts have been found for GENSENG.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at PRECISE-TBI .

Thangam M, et al. (2015) CRCDA--Comprehensive resources for cancer NGS data analysis. Database : the journal of biological databases and curation, 2015.