

Resource Summary Report

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

[CGARS](#)

RRID:SCR_006404

Type: Tool

Proper Citation

CGARS (RRID:SCR_006404)

Resource Information

URL: <http://www.uni-koeln.de/med-fak/cgars/>

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Description: Software package to dissect random from non-random patterns in copy number data and thereby to assess significantly enriched somatic copy number aberrations (SCNA) across a set of tumor specimens or cell lines.

Abbreviations: CGARS

Synonyms: CGARS: Cancer Genome Analysis by Rank Sums, Cancer Genome Analysis by Rank Sums

Resource Type: software resource

Defining Citation: [PMID:24413525](#)

Keywords: genome, analysis

Related Condition: Cancer

Availability: GNU General Public License, v3 or later

Resource Name: CGARS

Resource ID: SCR_006404

Alternate IDs: OMICS_02210

Record Creation Time: 20220129T080236+0000

Record Last Update: 20260905T132553+0000

Ratings and Alerts

No rating or validation information has been found for CGARS.

No alerts have been found for CGARS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.