

Resource Summary Report

Generated by [T1D](#) on Aug 1, 2026

[snapCGH](#)

RRID:SCR_012947

Type: Tool

Proper Citation

snapCGH (RRID:SCR_012947)

Resource Information

URL: <http://www.bioconductor.org/packages/2.12/bioc/html/snapCGH.html>

Proper Citation: snapCGH (RRID:SCR_012947)

Description: Software providing methods for segmenting, normalising and processing aCGH data; including plotting functions for visualising raw and segmented data for individual and multiple arrays.

Abbreviations: snapCGH

Resource Type: software resource

Resource Name: snapCGH

Resource ID: SCR_012947

Alternate IDs: OMICS_00734

Record Creation Time: 20220129T080313+0000

Record Last Update: 20260725T190744+0000

Ratings and Alerts

No rating or validation information has been found for snapCGH.

No alerts have been found for snapCGH.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Altmayer NC, et al. (2018) Gene amplification in mesenchymal stem cells and during differentiation towards adipocytes or osteoblasts. *Oncotarget*, 9(2), 1803.

La Fortezza M, et al. (2018) DamID profiling of dynamic Polycomb-binding sites in *Drosophila* imaginal disc development and tumorigenesis. *Epigenetics & chromatin*, 11(1), 27.

Chi C, et al. (2016) A Novel Graph-based Algorithm to Infer Recurrent Copy Number Variations in Cancer. *Cancer informatics*, 15(Suppl 2), 43.

Roessler S, et al. (2015) Integrative genomic and transcriptomic characterization of matched primary and metastatic liver and colorectal carcinoma. *International journal of biological sciences*, 11(1), 88.

Dib A, et al. (2009) A der(8)t(8;11) chromosome in the Karpas-620 myeloma cell line expresses only cyclin D1: yet both cyclin D1 and MYC are repositioned in close proximity to the 3'IGH enhancer. *DNA repair*, 8(3), 330.