## **Resource Summary Report**

Generated by NIF on Apr 29, 2025

# **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

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**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

**Funding:** 

Resource Name: snapCGH

Resource ID: SCR\_012947

Alternate IDs: OMICS\_00734

**Record Creation Time: 20220129T080313+0000** 

Record Last Update: 20250420T014625+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 NIF.

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.