Resource Summary Report

Generated by dkNET on Apr 26, 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

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

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 dkNET.

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.