

Resource Summary Report

Generated by [nidm-terms](#) on Sep 1, 2026

SnoopCGH

RRID:SCR_004420

Type: Tool

Proper Citation

SnoopCGH (RRID:SCR_004420)

Resource Information

URL: <http://snoopcgh.sourceforge.net/>

Proper Citation: SnoopCGH (RRID:SCR_004420)

Description: A java desktop application for visualising and exploring comparative genomic hybridization (CGH) data.

Abbreviations: SnoopCGH

Resource Type: software resource

Defining Citation: [PMID:19687029](#)

Keywords: bio.tools

Resource Name: SnoopCGH

Resource ID: SCR_004420

Alternate IDs: biotools:snoopcgh, OMICS_00736

Alternate URLs: <https://bio.tools/snoopcgh>

Record Creation Time: 20220129T080224+0000

Record Last Update: 20260829T182207+0000

Ratings and Alerts

No rating or validation information has been found for SnoopCGH.

No alerts have been found for SnoopCGH.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Farslow JC, et al. (2015) Rapid Increase in frequency of gene copy-number variants during experimental evolution in *Caenorhabditis elegans*. BMC genomics, 16, 1044.

Karimpour-Fard A, et al. (2010) A survey of analysis software for array-comparative genomic hybridisation studies to detect copy number variation. Human genomics, 4(6), 421.