

# Resource Summary Report

Generated by [RRID](#) on Aug 6, 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:** 20260801T190241+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 [RRID](#) .

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.