

Resource Summary Report

Generated by [NIF](#) on Aug 7, 2026

[cghMCR](#)

RRID:SCR_012898

Type: Tool

Proper Citation

cghMCR (RRID:SCR_012898)

Resource Information

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

Proper Citation: cghMCR (RRID:SCR_012898)

Description: Software package that provides functions to identify genomic regions of interest based on segmented copy number data from multiple samples.

Abbreviations: cghMCR

Resource Type: software resource

Resource Name: cghMCR

Resource ID: SCR_012898

Alternate IDs: OMICS_00710

Record Creation Time: 20220129T080313+0000

Record Last Update: 20260801T190453+0000

Ratings and Alerts

No rating or validation information has been found for cghMCR.

No alerts have been found for cghMCR.

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](#) .

Audrito V, et al. (2022) Tumors carrying BRAF-mutations over-express NAMPT that is genetically amplified and possesses oncogenic properties. Journal of translational medicine, 20(1), 118.

Rose Li Y, et al. (2020) Mutational signatures in tumours induced by high and low energy radiation in Trp53 deficient mice. Nature communications, 11(1), 394.

Iyer P, et al. (2019) ERBB2 and KRAS alterations mediate response to EGFR inhibitors in early stage gallbladder cancer. International journal of cancer, 144(8), 2008.

Zhang W, et al. (2019) Mutational signatures and the genomic landscape of betel quid chewing-associated tongue carcinoma. Cancer medicine, 8(2), 701.

Zhang K, et al. (2017) Screening of candidate key genes associated with human osteosarcoma using bioinformatics analysis. Oncology letters, 14(3), 2887.