

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

ADaCGH2

RRID:SCR_001981

Type: Tool

Proper Citation

ADaCGH2 (RRID:SCR_001981)

Resource Information

URL: <http://www.bioconductor.org/packages/release/bioc/html/ADaCGH2.html>

Proper Citation: ADaCGH2 (RRID:SCR_001981)

Description: Software for analysis and plotting of array comparative genomic hybridization (CGH) data. It allows usage of Circular Binary Segmentation, wavelet-based smoothing (both as in Liu et al., and HaarSeg as in Ben-Yaacov and Eldar), HMM, BioHMM, GLAD, CGHseg. Most computations are parallelized (either via forking or with clusters, including MPI and sockets clusters) and use ff for storing data.

Synonyms: ADaCGH2 - Analysis of big data from aCGH experiments using parallel computing and ff objects

Resource Type: software resource

Defining Citation: [PMID:24532724](#)

Keywords: standalone software, mac os x, unix/linux, windows, r, copy number variant, microarray, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: ADaCGH2

Resource ID: SCR_001981

Alternate IDs: biotools:adacgh2, OMICS_03697

Alternate URLs: <https://bio.tools/adacgh2>

License: GNU General Public License, v3

Record Creation Time: 20220129T080210+0000

Record Last Update: 20260725T190517+0000

Ratings and Alerts

No rating or validation information has been found for ADaCGH2.

No alerts have been found for ADaCGH2.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.