

Resource Summary Report

Generated by [NIF](#) on Aug 26, 2026

CNTools

RRID:SCR_000281

Type: Tool

Proper Citation

CNTools (RRID:SCR_000281)

Resource Information

URL: <http://www.bioconductor.org/packages/devel/bioc/html/CNTools.html>

Proper Citation: CNTools (RRID:SCR_000281)

Description: Software package that provides tools to convert the output of segmentation analysis using DNACopy to a matrix structure with overlapping segments as rows and samples as columns so that other computational analyses can be applied to segmented data.

Abbreviations: CNTools

Synonyms: CNTools - Convert segment data into a region by sample matrix to allow for other high level computational analyses

Resource Type: software resource

Keywords: copy number variation, microarray

Availability: Free, Available for download, Freely available

Resource Name: CNTools

Resource ID: SCR_000281

Alternate IDs: OMICS_02061

License: GNU Lesser General Public License

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260815T182146+0000

Ratings and Alerts

No rating or validation information has been found for CNTools.

No alerts have been found for CNTools.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Michal M, et al. (2026) FET-Rearranged Myoepithelial Tumors Are Clinically Heterogeneous and Epigenetically Distinct from PLAG1-Rearranged Adnexal and Salivary Gland Myoepithelial Tumors. Clinical cancer research : an official journal of the American Association for Cancer Research, 32(3), 628.