Resource Summary Report

Generated by nidm-terms on Jul 28, 2026

Computational Cancer Genomics Group

RRID:SCR_000772
Type: Tool

Proper Citation

Computational Cancer Genomics Group (RRID:SCR_000772)

Resource Information

URL: http://www.isrec.isb-sib.ch/

Proper Citation: Computational Cancer Genomics Group (RRID:SCR_000772)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 23,2022. The Computational Cancer Genomics (CCG) group is dedicated to the development of analysis tools and databases relating molecular sequences and biological functions. Sponsors: This group is supported by the Swiss Institute of Bioinformatics (SIB).

Synonyms: CCG

Resource Type: database, data analysis software, software resource, software application, data processing software, data or information resource

Keywords: eukaryotic, expression, function, gene, analyzer, annotation, biological, clustering, computational, data, genome, in vitro, mapping, messengerrna, molecular, mpss, mrna, one-dimensional, organism, promoter, sage, sequence, snp, software, tag, technology, tool, transcription, transcriptome

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Computational Cancer Genomics Group

Resource ID: SCR_000772

Alternate IDs: nif-0000-25561

Record Creation Time: 20220129T080203+0000

Record Last Update: 20260728T043102+0000

Ratings and Alerts

No rating or validation information has been found for Computational Cancer Genomics Group.

No alerts have been found for Computational Cancer Genomics Group.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at nidm-terms .

Shao G, et al. (2015) A Combinational Clustering Based Method for cDNA Microarray Image Segmentation. PloS one, 10(8), e0133025.

Morozov SY, et al. (2014) Plant 4/1 protein: potential player in intracellular, cell-to-cell and long-distance signaling. Frontiers in plant science, 5, 26.

Lin CH, et al. (2001) A small domain of CBP/p300 binds diverse proteins: solution structure and functional studies. Molecular cell, 8(3), 581.