

Resource Summary Report

Generated by [T1D](#) on Sep 19, 2026

[CGHweb](#)

RRID:SCR_010923

Type: Tool

Proper Citation

CGHweb (RRID:SCR_010923)

Resource Information

URL: <http://compbio.med.harvard.edu/CGHweb/>

Proper Citation: CGHweb (RRID:SCR_010923)

Description: Data analysis service enabling users to analyse their array-CGH data with multiple algorithms simultaneously.

Abbreviations: CGHweb

Resource Type: analysis service resource, data analysis service, production service resource, service resource

Resource Name: CGHweb

Resource ID: SCR_010923

Alternate IDs: OMICS_00713

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260919T195725+0000

Ratings and Alerts

No rating or validation information has been found for CGHweb.

No alerts have been found for CGHweb.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 12 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Przybyl J, et al. (2022) Detection of MDM2 amplification by shallow whole genome sequencing of cell-free DNA of patients with dedifferentiated liposarcoma. *PloS one*, 17(1), e0262272.

Montaño A, et al. (2020) Comprehensive Custom NGS Panel Validation for the Improvement of the Stratification of B-Acute Lymphoblastic Leukemia Patients. *Journal of personalized medicine*, 10(3).

Thybert D, et al. (2018) Repeat associated mechanisms of genome evolution and function revealed by the *Mus caroli* and *Mus pahari* genomes. *Genome research*, 28(4), 448.

Breman AM, et al. (2016) Evidence for feasibility of fetal trophoblastic cell-based noninvasive prenatal testing. *Prenatal diagnosis*, 36(11), 1009.

Forero-Castro M, et al. (2016) Genome-Wide DNA Copy Number Analysis of Acute Lymphoblastic Leukemia Identifies New Genetic Markers Associated with Clinical Outcome. *PloS one*, 11(2), e0148972.

Jacobs K, et al. (2016) Higher-Density Culture in Human Embryonic Stem Cells Results in DNA Damage and Genome Instability. *Stem cell reports*, 6(3), 330.

Merrick CJ, et al. (2015) Functional analysis of sirtuin genes in multiple *Plasmodium falciparum* strains. *PloS one*, 10(3), e0118865.

Ryba T, et al. (2012) Abnormal developmental control of replication-timing domains in pediatric acute lymphoblastic leukemia. *Genome research*, 22(10), 1833.

Lai LA, et al. (2012) Pan-colonic field defects are detected by CGH in the colons of UC patients with dysplasia/cancer. *Cancer letters*, 320(2), 180.

Ren M, et al. (2012) Acute progression of BCR-FGFR1 induced murine B-lympho/myeloproliferative disorder suggests involvement of lineages at the pro-B cell stage. *PloS one*, 7(6), e38265.

Fontanesi L, et al. (2012) Exploring copy number variation in the rabbit (*Oryctolagus cuniculus*) genome by array comparative genome hybridization. *Genomics*, 100(4), 245.

Fontanesi L, et al. (2011) A first comparative map of copy number variations in the sheep genome. *Genomics*, 97(3), 158.