

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 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, service resource, data analysis service, production service resource

Resource Name: CGHweb

Resource ID: SCR_010923

Alternate IDs: OMICS_00713

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260728T043345+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 [nidm-terms](#) .

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.

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

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