# **Resource Summary Report**

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# **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:** data analysis service, service resource, production service resource, analysis service resource

Funding:

Resource Name: CGHweb

Resource ID: SCR\_010923

Alternate IDs: OMICS\_00713

Record Creation Time: 20220129T080301+0000

Record Last Update: 20250524T060419+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 NIF.

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 Blympho/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.