## **Resource Summary Report**

Generated by RRID on Apr 19, 2025

# **CGH Explorer**

RRID:SCR\_010920

Type: Tool

### **Proper Citation**

CGH Explorer (RRID:SCR\_010920)

#### **Resource Information**

**URL:** http://www.softgenetics.com/CGHExplorer.html

**Proper Citation:** CGH Explorer (RRID:SCR\_010920)

**Description:** An easy-to-use software tool for analyzing two color copy number alteration arrays from multiple platforms, including Agilent Technologies, Illumina, AffyMetrix, NimbleGen and others.

Abbreviations: CGH Explorer

**Resource Type:** software resource

**Funding:** 

Availability: Commercially available

**Resource Name:** CGH Explorer

Resource ID: SCR\_010920

Alternate IDs: OMICS\_00708

**Record Creation Time:** 20220129T080301+0000

Record Last Update: 20250410T070030+0000

## Ratings and Alerts

No rating or validation information has been found for CGH Explorer.

No alerts have been found for CGH Explorer.

#### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at RRID.

, et al. (2018) Assessment of established techniques to determine developmental and malignant potential of human pluripotent stem cells. Nature communications, 9(1), 1925.

Davidsson J, et al. (2008) Deletion of the SCN gene cluster on 2q24.4 is associated with severe epilepsy: an array-based genotype-phenotype correlation and a comprehensive review of previously published cases. Epilepsy research, 81(1), 69.