Resource Summary Report

Generated by kravitz2 on Sep 18, 2026

CytoSure Interpret Software

RRID:SCR_010926
Type: Tool

Proper Citation

CytoSure Interpret Software (RRID:SCR_010926)

Resource Information

URL: http://www.ogt.co.uk/products/246_cytosure_interpret_software

Proper Citation: CytoSure Interpret Software (RRID:SCR_010926)

Description: A powerful and easy-to-use software package for the analysis of aCGH data, offering an impressive combination of features.

Abbreviations: CytoSure Interpret Software

Resource Type: commercial organization, software resource

Availability: Commercial license

Resource Name: CytoSure Interpret Software

Resource ID: SCR_010926

Alternate IDs: OMICS_00718

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260912T200331+0000

Ratings and Alerts

No rating or validation information has been found for CytoSure Interpret Software.

No alerts have been found for CytoSure Interpret Software.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 10 mentions in open access literature.

Listed below are recent publications. The full list is available at kravitz2 .

Madritsch S, et al. (2024) Aneuploidy detection in pooled polar bodies using rapid nanopore sequencing. Journal of assisted reproduction and genetics, 41(5), 1261.

Kinnart I, et al. (2024) Elevated ?-synuclein levels inhibit mitophagic flux. NPJ Parkinson's disease, 10(1), 80.

Nordenskjöld A, et al. (2023) Copy number variants suggest different molecular pathways for the pathogenesis of bladder exstrophy. American journal of medical genetics. Part A, 191(2), 378.

Dell'Edera D, et al. (2021) Mayer-Rokitansky-Küster-Hauser syndrome with 22q11.21 microduplication: a case report. Journal of medical case reports, 15(1), 208.

Kutkowska-Ka?mierczak A, et al. (2021) Wide Fontanels, Delayed Speech Development and Hoarse Voice as Useful Signs in the Diagnosis of KBG Syndrome: A Clinical Description of 23 Cases with Pathogenic Variants Involving the ANKRD11 Gene or Submicroscopic Chromosomal Rearrangements of 16q24.3. Genes, 12(8).

Colley E, et al. (2020) Cell-Free DNA in the Investigation of Miscarriage. Journal of clinical medicine, 9(11).

Lindstrand A, et al. (2019) From cytogenetics to cytogenomics: whole-genome sequencing as a first-line test comprehensively captures the diverse spectrum of disease-causing genetic variation underlying intellectual disability. Genome medicine, 11(1), 68.

Vincent-Chong VK, et al. (2017) Genome wide profiling in oral squamous cell carcinoma identifies a four genetic marker signature of prognostic significance. PloS one, 12(4), e0174865.

Barnes DJ, et al. (2016) A germline mutation of CDKN2A and a novel RPLP1-C19MC fusion detected in a rare melanotic neuroectodermal tumor of infancy: a case report. BMC cancer, 16, 629.

Baskin B, et al. (2014) Complex genomic rearrangements in the dystrophin gene due to replication-based mechanisms. Molecular genetics & genomic medicine, 2(6), 539.