Resource Summary Report

Generated by NIF on Apr 18, 2025

JISTIC

RRID:SCR_003482
Type: Tool

Proper Citation

JISTIC (RRID:SCR_003482)

Resource Information

URL: http://www.c2b2.columbia.edu/danapeerlab/html/jistic.html

Proper Citation: JISTIC (RRID:SCR_003482)

Description: Software tool for analyzing datasets of genome-wide copy number variation to identify driver aberrations in cancer.

Abbreviations: JISTIC

Resource Type: software resource

Defining Citation: PMID:20398270

Keywords: copy number variation, candidate gene, gene

Related Condition: Cancer

Funding:

Availability: Free, Public

Resource Name: JISTIC

Resource ID: SCR_003482

Alternate IDs: OMICS_02297

Record Creation Time: 20220129T080219+0000

Record Last Update: 20250410T065010+0000

Ratings and Alerts

No rating or validation information has been found for JISTIC.

No alerts have been found for JISTIC.

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 NIF.

Guo J, et al. (2018) Germline and somatic variations influence the somatic mutational signatures of esophageal squamous cell carcinomas in a Chinese population. BMC genomics, 19(1), 538.

Johnson N, et al. (2015) SubPatCNV: approximate subspace pattern mining for mapping copy-number variations. BMC bioinformatics, 16, 16.