Resource Summary Report

Generated by Kravitz on Sep 10, 2026

infercnvpy

RRID:SCR_025804
Type: Tool

Proper Citation

infercnvpy (RRID:SCR_025804)

Resource Information

URL: https://holonet-tutorial.readthedocs.io/en/latest/index.html

Proper Citation: infercnvpy (RRID:SCR_025804)

Description: Software application to infer copy number variation from single-cell transcriptomics data. Scalable Python library to infer CNV events from single cell transcriptomics data.

Resource Type: software library, software resource, software toolkit

Keywords: infer copy number variation, copy number variation, Copy Number Variation, CNV, single-cell transcriptomics data,

Availability: Free, Freely available,

Resource Name: infercnvpy

Resource ID: SCR_025804

Record Creation Time: 20240927T053302+0000

Record Last Update: 20260905T133507+0000

Ratings and Alerts

No rating or validation information has been found for infercnvpy.

No alerts have been found for infercnvpy.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Liu Y, et al. (2025) SpatialSNV: A novel method for identifying and analyzing spatially resolved SNVs in tumor microenvironments. GigaScience, 14.

Lin GT, et al. (2025) Combining Apatinib and Oxaliplatin Remodels the Immunosuppressive Tumor Microenvironment and Sensitizes Desert-Type Gastric Cancer to Immunotherapy. Cancer research, 85(11), 2117.

Strati P, et al. (2025) A Phase I Trial of Evorpacept, Lenalidomide, and Rituximab for Patients with B-Cell Non-Hodgkin Lymphoma. Clinical cancer research : an official journal of the American Association for Cancer Research, 31(19), 4079.