Resource Summary Report

Generated by RRID on Aug 2, 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 resource, software library, 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: 20260801T191339+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 2 mentions in open access literature.

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

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.

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