Resource Summary Report

Generated by NIF on Apr 20, 2025

<u>qips</u>

RRID:SCR_001092 Type: Tool

Proper Citation

qips (RRID:SCR_001092)

Resource Information

URL: http://sourceforge.net/projects/qips/

Proper Citation: qips (RRID:SCR_001092)

Description: A software package for analyzing ChIP-seq (Chromatin ImmunoPrecipitation on sequencing) data that finds enriched regions of arbitrary lengths and is therefore especially suited for analyzing ChIP-seq of histone marks or polymerase.

Abbreviations: qips

Resource Type: software resource

Keywords: command-line, c++, python

Funding:

Availability: Public Domain

Resource Name: gips

Resource ID: SCR_001092

Alternate IDs: OMICS_00457

Record Creation Time: 20220129T080205+0000

Record Last Update: 20250420T014021+0000

Ratings and Alerts

No rating or validation information has been found for qips.

No alerts have been found for qips.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.