

Resource Summary Report

Generated by PRECISE-TBI on Jul 31, 2026

qips

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

Availability: Free, Available for download, Freely available

Resource Name: qips

Resource ID: SCR_001092

Alternate IDs: OMICS_00457

Record Creation Time: 20220129T080205+0000

Record Last Update: 20260725T190456+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.