## **Resource Summary Report**

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## <u>iPiG</u>

RRID:SCR 016164

Type: Tool

### **Proper Citation**

iPiG (RRID:SCR\_016164)

#### Resource Information

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

Proper Citation: iPiG (RRID:SCR\_016164)

**Description:** Standalone software tool for the integration of peptide identifications from mass

spectrometry experiments into existing genome browser visualizations.

Abbreviations: iPiG

Synonyms: iPiG: Integrating Peptide Spectrum Matches Into Genome Browser

Visualizations

**Resource Type:** standalone software, software resource, software application

Defining Citation: PMID:23226516, DOI:10.1371/journal.pone.0050246

**Keywords:** integration, peptide, spectrum, match, genome, browser, visualization,

experiment, pms, ms, bio.tools

Funding: Robert Koch-Institute

Availability: Free, Available for download

Resource Name: iPiG

Resource ID: SCR\_016164

Alternate IDs: biotools:ipig, OMICS\_06913

Alternate URLs: https://bio.tools/ipig, https://sources.debian.org/src/ipig/

License: BSD License

**Record Creation Time:** 20220129T080329+0000

Record Last Update: 20250417T065542+0000

### **Ratings and Alerts**

No rating or validation information has been found for iPiG.

No alerts have been found for iPiG.

### **Data and Source Information**

Source: SciCrunch Registry

# Usage and Citation Metrics

We have not found any literature mentions for this resource.