

# Resource Summary Report

Generated by [NIF](#) on Sep 3, 2026

## iPiG

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:** software application, software resource, standalone software

**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:** 20260904T000456+0000

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## Ratings and Alerts

No rating or validation information has been found for iPiG.

No alerts have been found for iPiG.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.