

# Resource Summary Report

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## strap-base

RRID:SCR\_024351

Type: Tool

### Proper Citation

strap-base (RRID:SCR\_024351)

### Resource Information

**URL:** <http://www.bioinformatics.org/strap/>

**Proper Citation:** strap-base (RRID:SCR\_024351)

**Description:** Software tool as Intuitive Editor for annotated multiple Sequence and Structure Alignments.

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

**Keywords:** editor for annotated multiple Sequence and Structure Alignments,

**Availability:** Free, Available for download, Freely available,

**Resource Name:** strap-base

**Resource ID:** SCR\_024351

**Old URLs:** <https://sources.debian.org/src/strap-base/>

**Record Creation Time:** 20230830T050217+0000

**Record Last Update:** 20260801T191120+0000

### Ratings and Alerts

No rating or validation information has been found for strap-base.

No alerts have been found for strap-base.

### Data and Source Information

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

## Usage and Citation Metrics

We have not found any literature mentions for this resource.