

Resource Summary Report

Generated by [RRID](#) on Sep 18, 2026

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 application, software resource

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: 20260912T200308+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 found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Zhao J, et al. (2026) Novel compound heterozygous PKHD1 mutations in a Chinese ARPKD pedigree and analysis of genotype-phenotype correlations. *Frontiers in medicine*, 13, 1741795.

Duan Y, et al. (2023) Leader peptide removal in lasso peptide biosynthesis based on penultimate isoleucine residue. *Frontiers in microbiology*, 14, 1181125.

Tetreau G, et al. (2022) De novo determination of mosquitocidal Cry11Aa and Cry11Ba structures from naturally-occurring nanocrystals. *Nature communications*, 13(1), 4376.