

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 7, 2026

CoverageCalculator

RRID:SCR_005352

Type: Tool

Proper Citation

CoverageCalculator (RRID:SCR_005352)

Resource Information

URL: <http://covcal.sourceforge.net/>

Proper Citation: CoverageCalculator (RRID:SCR_005352)

Description: Small and very fast utility to calculate X-coverage from Next-Generation-Sequencing data.

Abbreviations: CoverageCalculator

Resource Type: software resource

Keywords: next-generation sequencing

Resource Name: CoverageCalculator

Resource ID: SCR_005352

Alternate IDs: OMICS_01164

Record Creation Time: 20220129T080229+0000

Record Last Update: 20260905T132534+0000

Ratings and Alerts

No rating or validation information has been found for CoverageCalculator.

No alerts have been found for CoverageCalculator.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Trivedi R, et al. (2019) Amino acid substitution scoring matrices specific to intrinsically disordered regions in proteins. Scientific reports, 9(1), 16380.

Hess M, et al. (2016) Addressing inaccuracies in BLOSUM computation improves homology search performance. BMC bioinformatics, 17, 189.