

Resource Summary Report

Generated by [NIF](#) on Aug 20, 2026

Crossbow

RRID:SCR_013306

Type: Tool

Proper Citation

Crossbow (RRID:SCR_013306)

Resource Information

URL: <http://bowtie-bio.sourceforge.net/crossbow/index.shtml>

Proper Citation: Crossbow (RRID:SCR_013306)

Description: A scalable software pipeline for whole genome resequencing analysis.

Abbreviations: Crossbow

Resource Type: software resource

Defining Citation: [PMID:19930550](#), [DOI:10.1186/gb-2009-10-11-r134](#)

Keywords: mapreduce/hadoop, bio.tools

Resource Name: Crossbow

Resource ID: SCR_013306

Alternate IDs: biotools:crossbow, OMICS_00284

Alternate URLs: <https://bio.tools/crossbow>

Old URLs: <https://sources.debian.org/src/crossbow/>

Record Creation Time: 20220129T080315+0000

Record Last Update: 20260815T182455+0000

Ratings and Alerts

No rating or validation information has been found for Crossbow.

No alerts have been found for Crossbow.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Amberg KL, et al. (2025) Proteasome regulation of petite-negativity in fission yeast. BMC biology, 23(1), 302.

Amberg KL, et al. (2024) Proteasome regulation of petite-negativity in fission yeast. bioRxiv : the preprint server for biology.

Lelieveld SH, et al. (2016) Novel bioinformatic developments for exome sequencing. Human genetics, 135(6), 603.

Testa A, et al. (2016) A Formal Methodology to Design and Deploy Dependable Wireless Sensor Networks. Sensors (Basel, Switzerland), 17(1).

Xia F, et al. (2007) Wireless Sensor/Actuator Network Design for Mobile Control Applications. Sensors (Basel, Switzerland), 7(10), 2157.