

# Resource Summary Report

Generated by [ASWG](#) on Aug 3, 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:** 20260801T190448+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](#)

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## Usage and Citation Metrics

We found 5 mentions in open access literature.

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

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.

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

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

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