## **Resource Summary Report**

Generated by <u>dkNET</u> on May 6, 2025

# 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

**Funding:** 

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: 20250420T014640+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: <u>SciCrunch Registry</u>

### **Usage and Citation Metrics**

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>dkNET</u>.

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.