Resource Summary Report

Generated by NIF on Apr 18, 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: 20250410T070352+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 4 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

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.